SUPPLEMENTAL TABLE 3: Part 1

			Summary Difference ALLC-JTO					
Uniprot ID	Gene name	Description	Average	St. Dev.	p-value	q-value (Stor	High-confidence Int	Secretory
XXXXXX	LC	lambda LC constant region (JTO-ALLC consensus)						
P23284	PPIB	Peptidyl-prolyl cis-trans isomerase B OS	0.69542552	0.05626755	0.00217508	0.02457032	TRUE	TRUE
Q9BS26	ERP44	Endoplasmic reticulum resident protein	0.95671444	0.11646843	0.00490373	0.04154552	TRUE	TRUE
P40227	CCT6A	T-complex protein 1 subunit zeta OS	0.3154045	0.004527	0.0064609	0.04691847	TRUE	FALSE
P11021	HSPA5	78 kDa glucose-regulated protein OS	0.68079165	0.10816554	0.00830976	0.04693475	TRUE	TRUE
P13667	PDIA4	Protein disulfide-isomerase A4 OS	0.53287846	0.09201742	0.00979368	0.04329089	TRUE	TRUE
P34931	HSPA1L	Heat shock 70 kDa protein 1-like OS	0.49783458	0.08995893	0.01070968	0.04032658	TRUE	FALSE
P53396	ACLY	ATP-citrate synthase OS	0.56025438	0.10372476	0.01123329	0.03938107	TRUE	FALSE
P30040	ERP29	Endoplasmic reticulum resident protein	0.54735697	0.1025182	0.01149218	0.0376894	TRUE	TRUE
Q02809	PLOD1	Procollagen-lysine,2-oxoglutarate 5-dio	0.5176443	0.09739302	0.01159489	0.03683793	TRUE	TRUE
P04843	RPN1	Dolichyl-diphosphooligosaccharide--pro	0.46190257	0.10911944	0.01809945	0.03833564	TRUE	TRUE
P49458	SRP9	Signal recognition particle 9 kDa protei	0.30585448	0.0756927	0.01981072	0.03800169	TRUE	FALSE
O76094	SRP72	Signal recognition particle 72 kDa prote	0.41628837	0.11036785	0.0226376	0.03712079	TRUE	TRUE
P54652	HSPA2	Heat shock-related 70 kDa protein 2 OS	0.4943038	0.14405172	0.02716114	0.03889271	TRUE	FALSE
Q8NBJ5	GLT25D1	Procollagen galactosyltransferase 1 OS	0.50942241	0.15233385	0.02853707	0.03920633	TRUE	TRUE
Q9UBS4	DNAJB11	DnaJ homolog subfamily B member 11	0.64720721	0.1935587	0.02854348	0.03869228	TRUE	TRUE
Q9NYU2	UGGT1	UDP-glucose:glycoprotein glucosyltrans	0.53757228	0.16109045	0.02865238	0.03783107	TRUE	TRUE
Q07065	CKAP4	Cytoskeleton-associated protein 4 OS	0.58834699	0.18706834	0.03208559	0.03837688	TRUE	TRUE
Q14697	GANAB	Neutral alpha-glucosidase AB OS	0.47352992	0.16104364	0.03645874	0.03985633	TRUE	TRUE
Q13162	PRDX4	Peroxiredoxin-4 OS	0.65739471	0.23786583	0.04097689	0.03787258	TRUE	FALSE
Q8IXB1	DNAJC10	DnaJ homolog subfamily C member 10	0.66849959	0.24448309	0.0418075	0.03761442	TRUE	TRUE
P49257	LMAN1	Protein ERGIC-53 OS	0.43946572	0.16297254	0.0429122	0.03728838	TRUE	TRUE
Q9Y4L1	HYOU1	Hypoxia up-regulated protein 1 OS	0.54974645	0.20598154	0.0437482	0.03737592	TRUE	TRUE
Q969V3	NCLN	Nicalin OS	0.4665241	0.04707458	0.04534633	0.03841842	TRUE	TRUE
Q9Y2B0	CNPY2	Protein canopy homolog 2 OS	0.65923852	0.07022414	0.04786187	0.03956062	TRUE	TRUE
Q08380	LGALS3BP	Galectin-3-binding protein OS	0.49443467	0.19589289	0.04854518	0.03917005	TRUE	FALSE
P27797	CALR	Calreticulin OS	0.26880205	0.10744461	0.0493487	0.03859321	TRUE	TRUE
Q8NBS9	TXNDC5	Thioredoxin domain-containing protein	0.47390614	0.0546591	0.05180542	0.03872709	TRUE	TRUE
P14314	PRKCSH	Glucosidase 2 subunit beta OS	0.57531657	0.25720577	0.06062765	0.04136786	TRUE	TRUE
Q15293	RCN1	Reticulocalbin-1 OS	0.54223496	0.25206188	0.06507816	0.04161182	TRUE	TRUE
Q71U36	TUBA1A	Tubulin alpha-1A chain OS	0.64364149	0.096397	0.067169	0.04215339	TRUE	FALSE
P07237	P4HB	Protein disulfide-isomerase OS	0.57502063	0.27346323	0.06781071	0.04203713	TRUE	TRUE
P46977	STT3A	Dolichyl-diphosphooligosaccharide--pro	0.44289565	0.22105173	0.07394345	0.04320451	TRUE	TRUE
Q15084	PDIA6	Protein disulfide-isomerase A6 OS	0.48340395	0.24243649	0.07458217	0.04308251	TRUE	TRUE
O75976	CPD	Carboxypeptidase D OS	0.26571567	0.13387071	0.0751904	0.04318846	TRUE	FALSE
O43493	TGOLN2	Trans-Golgi network integral membran	-0.2776425	0.15422276	0.08928935	0.04631505	TRUE	TRUE
P55145	MANF	Mesencephalic astrocyte-derived neuro	0.57651093	0.12073615	0.09359444	0.04710611	TRUE	TRUE
P04844	RPN2	Dolichyl-diphosphooligosaccharide--pro	0.48457018	0.10303084	0.09500249	0.04757924	TRUE	TRUE
Q32P28	LEPRE1	Prolyl 3-hydroxylase 1 OS	0.57967243	0.3455253	0.10083658	0.04790523	TRUE	TRUE
O94905	ERLIN2	Erlin-2 OS	0.50210427	0.12573021	0.11156622	0.04785893	TRUE	TRUE
P49207	RPL34	60S ribosomal protein L34 OS	0.31803214	0.20167824	0.11197638	0.04763291	TRUE	FALSE
Q05707	COL14A1	Collagen alpha-1(XIV) chain OS	0.46959056	0.14152794	0.13367159	0.05070875	TRUE	TRUE
P68371	TUBB2C	Tubulin beta-2C chain OS	0.32894644	0.10046081	0.13539984	0.05042363	TRUE	FALSE
P30101	PDIA3	Protein disulfide-isomerase A3 OS	0.37229114	0.28246321	0.1499028	0.05237154	TRUE	TRUE
P13674	P4HA1	Prolyl 4-hydroxylase subunit alpha-1 OS	0.53102719	0.20338495	0.16837296	0.05539779	TRUE	TRUE
P23526	AHCY	Adenosylhomocysteinase OS	0.49125988	0.19435156	0.17365196	0.05640452	TRUE	FALSE
Q14204	DYNC1H1	Cytoplasmic dynein 1 heavy chain 1 OS	0.30762422	0.25831948	0.17523946	0.05637978	TRUE	FALSE
O60568	PLOD3	Procollagen-lysine,2-oxoglutarate 5-dio	0.50123062	0.2095614	0.18299585	0.05742154	TRUE	TRUE
O75718	CRTAP	Cartilage-associated protein OS	0.58237423	0.24494216	0.1840297	0.05756827	TRUE	TRUE
O14737	PDCD5	Programmed cell death protein 5 OS	0.47563335	0.42555015	0.19251348	0.05948998	TRUE	FALSE
P39656	DDOST	Dolichyl-diphosphooligosaccharide--pro	0.50685293	0.23051106	0.19807749	0.05993416	TRUE	TRUE
Q13217	DNAJC3	DnaJ homolog subfamily C member 3 O	0.57990915	0.26386161	0.19816571	0.05978293	TRUE	TRUE
Q96AY3	FKBP10	Peptidyl-prolyl cis-trans isomerase FKBI	0.51015838	0.23714534	0.20217215	0.06045344	TRUE	TRUE
O43405	COCH	Cochlin OS	0.27429859	0.25519994	0.2037021	0.06037817	TRUE	FALSE
Q07955	SRSF1	Serine/arginine-rich splicing factor 1 OS	0.73302468	0.36908041	0.21774765	0.06253581	TRUE	FALSE
Q9Y490	TLN1	Talin-1 OS	0.55728785	0.56570747	0.23007715	0.06391032	TRUE	FALSE
Q14257	RCN2	Reticulocalbin-2 OS	0.42473833	0.23154664	0.23422957	0.0645348	TRUE	TRUE
Q8NBJ4	GOLM1	Golgi membrane protein 1 OS	0.31282569	0.36464315	0.27563159	0.06818138	TRUE	TRUE
P14625	HSP90B1	Endoplasmin OS	0.25404979	0.30697454	0.28813723	0.06991397	TRUE	TRUE
P50454	SERPINH1	Serpin H1 OS	0.50094204	0.36222326	0.30089508	0.0714743	TRUE	TRUE
Q16706	MAN2A1	Alpha-mannosidase 2 OS	0.43410165	0.34016095	0.32211411	0.07375736	TRUE	TRUE
Q9NXG2	THUMPD1	THUMP domain-containing protein 1 O	0.28124672	0.38551124	0.33371626	0.07473088	TRUE	FALSE
P27824	CANX	Calnexin OS	0.18772137	0.26409254	0.34339028	0.07639244	TRUE	TRUE
Q96HE7	ERO1L	ERO1-like protein alpha OS	0.73193284	0.6700684	0.36574028	0.08013706	TRUE	TRUE
Q9H3N1	TMX1	Thioredoxin-related transmembrane pr	0.26952993	0.25421413	0.3744484	0.08169296	TRUE	TRUE
Q92896	GLG1	Golgi apparatus protein 1 OS	0.25910515	0.31919262	0.45620796	0.09582881	TRUE	TRUE
P02786	TFRC	Transferrin receptor protein 1 OS	0.11945296	0.34969495	0.61405168	0.1214564	TRUE	FALSE
O43852	CALU	Calumenin OS	0.13066568	0.3535585	0.69339942	0.1345336	TRUE	TRUE
Q00839	HNRNPU	Heterogeneous nuclear ribonucleopro	0.90094546	0.02323311	0.00022159	0.02252842	FALSE	FALSE
P35580	MYH10	Myosin-10 OS	0.47532959	0.01862016	0.00051112	0.02598192	FALSE	FALSE
Q13442	PDAP1	28 kDa heat- and acid-stable phosphop	0.68030042	0.02736822	0.00053904	0.01826738	FALSE	FALSE
P55209	NAP1L1	Nucleosome assembly protein 1-like 1 (	0.92425098	0.0436333	0.00074208	0.01886123	FALSE	FALSE
Q08211	DHX9	ATP-dependent RNA helicase A OS	0.40877765	0.00082356	0.00090693	0.01844083	FALSE	FALSE
P32969	RPL9	60S ribosomal protein L9 OS	0.64723747	0.04608152	0.00168541	0.02855836	FALSE	FALSE
P62917	RPL8	60S ribosomal protein L8 OS	0.43265561	0.03162291	0.00177599	0.02579409	FALSE	FALSE
Q99832	CCT7	T-complex protein 1 subunit eta OS	0.24784622	0.00108572	0.00197197	0.02506048	FALSE	FALSE
P34897	SHMT2	Serine hydroxymethyltransferase, mito	0.45880404	0.04283588	0.00289302	0.0294124	FALSE	FALSE
P46781	RPS9	40S ribosomal protein S9 OS	0.53446582	0.0513248	0.00305982	0.0282802	FALSE	FALSE
P62701	RPS4X	40S ribosomal protein S4, X isoform OS	0.35410228	0.04800535	0.0060706	0.04747523	FALSE	FALSE
P19338	NCL	Nucleolin OS	0.59862381	0.08413128	0.00651962	0.04418854	FALSE	FALSE
P35579	MYH9	Myosin-9 OS	0.42662055	0.06564281	0.00779948	0.04955917	FALSE	FALSE
O75822	EIF3J	Eukaryotic translation initiation factor 3	0.9477853	0.14905486	0.00814368	0.04870239	FALSE	FALSE
Q14566	MCM6	DNA replication licensing factor MCM6	0.3949525	0.06435769	0.00873515	0.0467407	FALSE	FALSE
P63261	ACTG1	Actin, cytoplasmic 2 OS	0.53793107	0.08774947	0.00875352	0.04449707	FALSE	FALSE
P07942	LAMB1	Laminin subunit beta-1 OS	0.35760703	0.00754598	0.00949822	0.04598345	FALSE	FALSE
P60866	RPS20	40S ribosomal protein S20 OS	0.5956753	0.10251426	0.00972869	0.04495832	FALSE	FALSE
P18621	RPL17	60S ribosomal protein L17 OS	0.40124888	0.06943287	0.00983418	0.04165866	FALSE	FALSE
P21333	FLNA	Filamin-A OS	0.56412778	0.09880971	0.01007216	0.04096013	FALSE	FALSE
Q02790	FKBP4	Peptidyl-prolyl cis-trans isomerase FKBI	0.81099561	0.14431165	0.01039045	0.04062933	FALSE	TRUE
P46782	RPS5	40S ribosomal protein S5 OS	0.50646198	0.09322811	0.01110698	0.04032892	FALSE	FALSE

P08670	VIM	Vimentin OS	0.83687458	0.15571418	0.01134423	0.03844432	FALSE	FALSE
P40429	RPL13A	60S ribosomal protein L13a OS	0.44921512	0.08538607	0.01182998	0.03644589	FALSE	FALSE
P62750	RPL23A	60S ribosomal protein L23a OS	0.51987582	0.10238514	0.01268321	0.0379253	FALSE	FALSE
Q9Y265	RUVBL1	RuvB-like 1 OS	0.45576504	0.0134848	0.01331696	0.03868261	FALSE	TRUE
P25205	MCM3	DNA replication licensing factor MCM3	0.56652806	0.11444698	0.01333188	0.03765022	FALSE	FALSE
P63220	RPS21	40S ribosomal protein S21 OS	0.63746361	0.12958902	0.01349714	0.03708674	FALSE	FALSE
P63241	EIF5A	Eukaryotic translation initiation factor 5	0.66797364	0.1392938	0.01418744	0.03795763	FALSE	TRUE
Q9HCN8	SDF2L1	Stromal cell-derived factor 2-like protei	0.6422463	0.0203561	0.01426545	0.0371877	FALSE	TRUE
P35268	RPL22	60S ribosomal protein L22 OS	0.57786846	0.124946	0.01522846	0.03870566	FALSE	FALSE
P61247	RPS3A	40S ribosomal protein S3a OS	0.47908854	0.1040168	0.01535192	0.03806777	FALSE	FALSE
P04181	OAT	Ornithine aminotransferase, mitochonc	0.54857317	0.11915589	0.01536528	0.03719372	FALSE	FALSE
P55010	EIF5	Eukaryotic translation initiation factor 5	0.75071052	0.16642075	0.01598947	0.03780457	FALSE	FALSE
Q9Y266	NUDC	Nuclear migration protein nudC OS	0.492985	0.11177298	0.01670683	0.03860291	FALSE	FALSE
P63104	YWHAZ	14-3-3 protein zeta/delta OS	0.56609811	0.12839127	0.01671739	0.03776891	FALSE	FALSE
Q9NSB2	KRT84	Keratin, type II cuticular Hb4 OS	0.48137746	0.01856051	0.01735248	0.0383515	FALSE	FALSE
P08107	HSPA1A	Heat shock 70 kDa protein 1A/1B OS	0.45054582	0.10446582	0.0174527	0.0377523	FALSE	FALSE
P09874	PARP1	Poly [ADP-ribose] polymerase 1 OS	0.41177633	0.09782809	0.01829925	0.03796783	FALSE	FALSE
P62249	RPS16	40S ribosomal protein S16 OS	0.33573533	0.0800885	0.01844498	0.03750479	FALSE	FALSE
P20290	BTF3	Transcription factor BTF3 OS	0.43835239	0.01889652	0.01939944	0.03867208	FALSE	FALSE
Q15691	MAPRE1	Microtubule-associated protein RP/EB f	0.46932145	0.02048144	0.01963891	0.03839659	FALSE	TRUE
P62937	PPIA	Peptidyl-prolyl cis-trans isomerase A OS	0.7827286	0.19692777	0.02045428	0.0385096	FALSE	FALSE
Q15366	PCBP2	Poly(rC)-binding protein 2 OS	0.62626551	0.1582719	0.02063307	0.03813993	FALSE	FALSE
P62241	RPS8	40S ribosomal protein S8 OS	0.59777636	0.1513416	0.0207045	0.03758853	FALSE	FALSE
P53618	COPB1	Coatomer subunit beta OS	0.38301144	0.01804081	0.02119576	0.0378053	FALSE	TRUE
O95433	AHSA1	Activator of 90 kDa heat shock protein	0.66787453	0.17231218	0.02147596	0.03764464	FALSE	TRUE
P31948	STIP1	Stress-induced-phosphoprotein 1 OS	0.65508918	0.17025679	0.02178278	0.03753531	FALSE	TRUE
Q5JTV8	TOR1AIP1	Torsin-1A-interacting protein 1 OS	0.59667327	0.15769304	0.02249976	0.03812459	FALSE	FALSE
Q7L1Q6	BZW1	Basic leucine zipper and W2 domain-co	0.734323	0.19424454	0.02253842	0.03756403	FALSE	FALSE
Q07020	RPL18	60S ribosomal protein L18 OS	0.43678284	0.1186278	0.02371664	0.03827288	FALSE	FALSE
Q0VDF9	HSPA14	Heat shock 70 kDa protein 14 OS	0.28507499	0.01584108	0.02500158	0.03971605	FALSE	FALSE
P11142	HSPA8	Heat shock cognate 71 kDa protein OS	0.48146284	0.13489463	0.02518213	0.03938743	FALSE	FALSE
P62753	RPS6	40S ribosomal protein S6 OS	0.62589814	0.17547189	0.02521251	0.03883745	FALSE	FALSE
Q99623	PHB2	Prohibitin-2 OS	0.32157002	0.09069414	0.02550466	0.0387011	FALSE	FALSE
P38646	HSPA9	Stress-70 protein, mitochondrial OS	0.52036813	0.15005585	0.02661643	0.03979417	FALSE	FALSE
P11717	IGF2R	Cation-independent mannose-6-phospl	0.2365008	0.01404104	0.02671018	0.03935559	FALSE	FALSE
P49736	MCM2	DNA replication licensing factor MCM2	0.59764005	0.17270733	0.02672599	0.03881632	FALSE	FALSE
P62269	RPS18	40S ribosomal protein S18 OS	0.32938824	0.0968064	0.02760528	0.03897967	FALSE	FALSE
P14618	PKM2	Pyruvate kinase isozymes M1/M2 OS	0.4842911	0.14291429	0.02782226	0.03874789	FALSE	FALSE
P05388	RPLP0	60S acidic ribosomal protein P0 OS	0.52486463	0.0333551	0.02858828	0.03824309	FALSE	FALSE
P62424	RPL7A	60S ribosomal protein L7a OS	0.56153627	0.03596168	0.02880916	0.03755041	FALSE	FALSE
P25789	PSMA4	Proteasome subunit alpha type-4 OS	0.24103739	0.01587694	0.02963015	0.03813163	FALSE	FALSE
Q13263	TRIM28	Transcription intermediary factor 1-bet	0.63249822	0.0421259	0.02995948	0.0380735	FALSE	FALSE
P04075	ALDOA	Fructose-bisphosphate aldolase A OS	0.50030381	0.15423225	0.03024831	0.03796599	FALSE	FALSE
P08621	SNRNP70	U1 small nuclear ribonucleoprotein 70	0.61653032	0.19224414	0.03091473	0.03832924	FALSE	FALSE
Q9UHB6	LIMA1	LIM domain and actin-binding protein 1	0.49621105	0.03522401	0.03192811	0.03910873	FALSE	FALSE
Q92616	GCN1L1	Translational activator GCN1 OS	0.23246807	0.01654989	0.03202068	0.03875519	FALSE	FALSE
P10809	HSPD1	60 kDa heat shock protein, mitochondr	0.42820976	0.13976554	0.03372508	0.0398688	FALSE	TRUE
P35749	MYH11	Myosin-11 OS	0.38719308	0.0290569	0.03375046	0.0394402	FALSE	FALSE
Q13151	HNRNPA0	Heterogeneous nuclear ribonucleopro	0.49045657	0.03707464	0.03399604	0.03927573	FALSE	FALSE
P09651	HNRNPA1	Heterogeneous nuclear ribonucleopro	0.58023143	0.19248743	0.03478173	0.03973194	FALSE	FALSE
P07195	LDHB	L-lactate dehydrogenase B chain OS	0.32776482	0.1090234	0.03495781	0.03948938	FALSE	FALSE
O00429	DNM1L	Dynamin-1-like protein OS	0.42555723	0.03394292	0.03586712	0.04007132	FALSE	TRUE
P51858	HDGF	Hepatoma-derived growth factor OS	0.95356962	0.32317956	0.03622049	0.04002627	FALSE	FALSE
P09012	SNRPA	U1 small nuclear ribonucleoprotein A O	0.61682871	0.21083739	0.03680765	0.03980969	FALSE	FALSE
Q7KZF4	SND1	Staphylococcal nuclease domain-contai	0.48612146	0.16681134	0.03708068	0.03968283	FALSE	FALSE
Q04917	YWHAH	14-3-3 protein eta OS	0.60458222	0.20748923	0.03709027	0.03927962	FALSE	FALSE
Q9NZI8	IGF2BP1	Insulin-like growth factor 2 mRNA-bind	0.29947492	0.0250957	0.03767875	0.03949147	FALSE	FALSE
P62979	RPS27A	Ubiquitin-40S ribosomal protein S27a C	0.52906622	0.18322768	0.03773157	0.0391433	FALSE	FALSE
Q99733	NAP1L4	Nucleosome assembly protein 1-like 4 (	0.62295795	0.05305018	0.03828857	0.03931991	FALSE	FALSE
P06733	ENO1	Alpha-enolase OS	0.43443182	0.15218204	0.03855369	0.03919625	FALSE	FALSE
Q9UNX3	RPL26L1	60S ribosomal protein L26-like 1 OS	0.35820773	0.12563527	0.03864332	0.0388984	FALSE	FALSE
Q04637	EIF4G1	Eukaryotic translation initiation factor 4	0.33945237	0.11916956	0.03871215	0.03858564	FALSE	FALSE
P61353	RPL27	60S ribosomal protein L27 OS	0.63313801	0.05455939	0.03874357	0.03824203	FALSE	FALSE
Q92499	DDX1	ATP-dependent RNA helicase DDX1 OS	0.54481831	0.04719574	0.03894699	0.03807318	FALSE	FALSE
P22102	GART	Trifunctional purine biosynthetic protei	0.61108842	0.0533529	0.03925259	0.03800647	FALSE	FALSE
P62316	SNRPD2	Small nuclear ribonucleoprotein Sm D2	0.63089057	0.05511508	0.03927625	0.03767061	FALSE	FALSE
P05387	RPLP2	60S acidic ribosomal protein P2 OS	0.99663697	0.35293371	0.03935056	0.03738916	FALSE	FALSE
P00338	LDHA	L-lactate dehydrogenase A chain OS	0.39110584	0.13857881	0.03939263	0.03708257	FALSE	FALSE
O43390	HNRNPR	Heterogeneous nuclear ribonucleopro	0.44671018	0.03927496	0.03952724	0.03686791	FALSE	TRUE
P46779	RPL28	60S ribosomal protein L28 OS	0.4421414	0.04051757	0.04119463	0.03773082	FALSE	FALSE
Q06830	PRDX1	Peroxiredoxin-1 OS	0.46522084	0.16949112	0.04150869	0.03767902	FALSE	FALSE
P26640	VARS	Valyl-tRNA synthetase OS	0.3668501	0.13476976	0.04216223	0.03760082	FALSE	FALSE
P41252	IARS	Isoleucyl-tRNA synthetase, cytoplasmic	0.60527191	0.2234486	0.04255045	0.03761707	FALSE	FALSE
P26373	RPL13	60S ribosomal protein L13 OS	0.31214282	0.11561114	0.0428118	0.03752184	FALSE	FALSE
Q14739	LBR	Lamin-B receptor OS	0.78219363	0.29276159	0.04366047	0.03761707	FALSE	FALSE
P62277	RPS13	40S ribosomal protein S13 OS	0.5541166	0.21533013	0.04682925	0.03934689	FALSE	FALSE
P22626	HNRNPA2B1	Heterogeneous nuclear ribonucleopro	0.61832234	0.06485109	0.04712737	0.0392728	FALSE	FALSE
P62318	SNRPD3	Small nuclear ribonucleoprotein Sm D3	0.68972755	0.07350384	0.04788258	0.03925857	FALSE	FALSE
P49327	FASN	Fatty acid synthase OS	0.38203262	0.15094531	0.04829866	0.03928291	FALSE	TRUE
Q16543	CDC37	Hsp90 co-chaperone Cdc37 OS	0.82701132	0.32932893	0.04900558	0.03923019	FALSE	FALSE
O14744	PRMT5	Protein arginine N-methyltransferase 5	0.42956157	0.17108176	0.04901811	0.03893365	FALSE	TRUE
Q9HB71	CACYBP	Calcyclin-binding protein OS	0.51200023	0.05590386	0.04905418	0.03866027	FALSE	FALSE
Q13765	NACA	Nascent polypeptide-associated compl	0.82890933	0.33160416	0.0494248	0.03835767	FALSE	FALSE
P25398	RPS12	40S ribosomal protein S12 OS	0.50354416	0.05682828	0.05069588	0.03904607	FALSE	FALSE
P53041	PPP5C	Serine/threonine-protein phosphatase	0.50397339	0.05690697	0.05072273	0.03877302	FALSE	FALSE
Q6P1N0	CC2D1A	Coiled-coil and C2 domain-containing p	0.70194673	0.08051193	0.0515195	0.03908818	FALSE	FALSE
Q86V81	THOC4	THO complex subunit 4 OS	0.7534925	0.08644202	0.05153012	0.03880663	FALSE	FALSE
P39023	RPL3	60S ribosomal protein L3 OS	0.36436651	0.14952595	0.05181119	0.0384487	FALSE	FALSE
Q8NE71	ABCF1	ATP-binding cassette sub-family F mem	1.05268568	0.12205668	0.0520784	0.03836694	FALSE	FALSE
P16615	ATP2A2	Sarcoplasmic/endoplasmic reticulum ca	0.52733484	0.21756729	0.05232679	0.03827259	FALSE	TRUE
P62851	RPS25	40S ribosomal protein S25 OS	0.2964736	0.12302619	0.05288648	0.03840565	FALSE	FALSE
Q9NSD9	FARSB	Phenylalanyl-tRNA synthetase beta cha	0.29214587	0.03477044	0.05345069	0.0385401	FALSE	FALSE
Q99613	EIF3C	Eukaryotic translation initiation factor 3	0.38914815	0.04657478	0.05374862	0.038482	FALSE	FALSE
P07437	TUBB	Tubulin beta chain OS	0.32342433	0.03910815	0.05430061	0.03860533	FALSE	FALSE

O75475	PSIP1	PC4 and SFRS1-interacting protein OS	0.57268649	0.24116185	0.05433718	0.03836306	FALSE	FALSE
P08133	ANXA6	Annexin A6 OS	0.47043619	0.05738995	0.05478057	0.03840936	FALSE	FALSE
P06454	PTMA	Prothymosin alpha OS	0.59016452	0.25910549	0.0586557	0.04084472	FALSE	FALSE
P62263	RPS14	40S ribosomal protein S14 OS	0.55937512	0.24863998	0.05999334	0.04149199	FALSE	FALSE
P08123	COL1A2	Collagen alpha-2(I) chain OS	1.4322414	0.19230845	0.06026253	0.04139656	FALSE	TRUE
P10599	TXN	Thioredoxin OS	0.63815308	0.28875658	0.06197185	0.04200314	FALSE	FALSE
P25705	ATP5A1	ATP synthase subunit alpha, mitochond	0.50344053	0.22946593	0.06279728	0.04228073	FALSE	FALSE
P27348	YWHAQ	14-3-3 protein theta OS	0.64769131	0.29557224	0.06293538	0.04209494	FALSE	FALSE
O43809	NUDT21	Cleavage and polyadenylation specificit	0.47735548	0.21880171	0.06344045	0.04215542	FALSE	FALSE
P62847	RPS24	40S ribosomal protein S24 OS	0.62138862	0.08789498	0.06346348	0.04189689	FALSE	FALSE
Q99615	DNAJC7	DnaJ homolog subfamily C member 7 O	0.59682959	0.27432389	0.06376029	0.04182127	FALSE	FALSE
O15372	EIF3H	Eukaryotic translation initiation factor 3	0.38033077	0.17617446	0.06466239	0.04214109	FALSE	FALSE
Q9UNE7	STUB1	E3 ubiquitin-protein ligase CHIP OS	0.61176056	0.08836296	0.06479635	0.04195942	FALSE	TRUE
Q9BQA1	WDR77	Methylosome protein 50 OS	0.41451932	0.05994498	0.06487333	0.04174339	FALSE	TRUE
P10515	DLAT	Dihydrolipoyllysine-residue acetyltransf	0.45773011	0.21526133	0.06645643	0.04222753	FALSE	FALSE
Q5VTE0	EEF1A1P5	Putative elongation factor 1-alpha-like	0.30819824	0.14549142	0.06691378	0.04225404	FALSE	FALSE
P52292	KPNA2	Importin subunit alpha-2 OS	0.49181923	0.0742275	0.06768365	0.04221577	FALSE	FALSE
P08579	SNRPB2	U2 small nuclear ribonucleoprotein B&i	0.51177037	0.07739563	0.06782021	0.04178821	FALSE	FALSE
P51572	BCAP31	B-cell receptor-associated protein 31 O	0.36536417	0.17710208	0.0701759	0.04297922	FALSE	TRUE
P50990	CCT8	T-complex protein 1 subunit theta OS	0.42682363	0.2069199	0.07019217	0.04273176	FALSE	FALSE
Q9P2E9	RRBP1	Ribosome-binding protein 1 OS	0.46228691	0.22449042	0.07040487	0.04260612	FALSE	TRUE
Q96A33	CCDC47	Coiled-coil domain-containing protein 4	0.65794164	0.10400952	0.07086825	0.04263277	FALSE	TRUE
O95373	IPO7	Importin-7 OS	0.63363697	0.10156205	0.07184669	0.04296714	FALSE	FALSE
Q9BXP5	SRRT	Serrate RNA effector molecule homolo	0.65537756	0.1050919	0.07187733	0.04273408	FALSE	FALSE
P49411	TUFM	Elongation factor Tu, mitochondrial OS	0.55233966	0.08857211	0.07187948	0.0424869	FALSE	FALSE
P08708	RPS17	40S ribosomal protein S17 OS	0.45931268	0.07522305	0.07339684	0.04313302	FALSE	FALSE
P55039	DRG2	Developmentally-regulated GTP-bindin	0.35371123	0.05852377	0.07414445	0.0430744	FALSE	FALSE
P60174	TPI1	Triosephosphate isomerase OS	0.73668339	0.3732207	0.07593838	0.04337304	FALSE	FALSE
P23396	RPS3	40S ribosomal protein S3 OS	0.76439986	0.39130459	0.07735171	0.04393347	FALSE	FALSE
P47813	EIF1AX	Eukaryotic translation initiation factor 1	0.80457652	0.41465545	0.07828094	0.04421423	FALSE	FALSE
Q9P258	RCC2	Protein RCC2 OS	0.73942948	0.13049631	0.07903642	0.0443943	FALSE	FALSE
P08238	HSP90AB1	Heat shock protein HSP 90-beta OS	0.82244971	0.42837583	0.07976025	0.04455472	FALSE	FALSE
Q9UNF0	PACSIN2	Protein kinase C and casein kinase subs	0.50697457	0.09144572	0.08076139	0.04486744	FALSE	FALSE
Q15424	SAFB	Scaffold attachment factor B1 OS	0.63416943	0.11561687	0.0816192	0.04509757	FALSE	FALSE
P49368	CCT3	T-complex protein 1 subunit gamma OS	0.421687	0.22363708	0.08233951	0.04524964	FALSE	FALSE
P62899	RPL31	60S ribosomal protein L31 OS	0.47947826	0.25477527	0.08261843	0.04515882	FALSE	FALSE
P83881	RPL36A	60S ribosomal protein L36a OS	0.48930334	0.09222855	0.08435295	0.04586034	FALSE	FALSE
P62854	RPS26	40S ribosomal protein S26 OS	0.85620412	0.46163266	0.08476051	0.0458368	FALSE	FALSE
P36578	RPL4	60S ribosomal protein L4 OS	0.58412679	0.11126686	0.08523499	0.04584951	FALSE	FALSE
Q15155	NOMO1	Nodal modulator 1 OS	0.46176899	0.08798231	0.0852567	0.04561981	FALSE	FALSE
P52597	HNRNPF	Heterogeneous nuclear ribonucleopro	0.57452742	0.10998175	0.08565315	0.04559199	FALSE	FALSE
P62258	YWHAE	14-3-3 protein epsilon OS	0.82578153	0.44860139	0.08588776	0.04547876	FALSE	FALSE
P0CG39	POTEJ	POTE ankyrin domain family member J	0.38611902	0.07470877	0.08656204	0.04559831	FALSE	FALSE
O15347	HMGB3	High mobility group protein B3 OS	0.76060128	0.41540851	0.08669479	0.04543284	FALSE	FALSE
P07900	HSP90AA1	Heat shock protein HSP 90-alpha OS	0.83306104	0.45926197	0.08812289	0.04594441	FALSE	FALSE
Q15365	PCBP1	Poly(rC)-binding protein 1 OS	0.70549937	0.39606032	0.09094885	0.04693638	FALSE	FALSE
P15880	RPS2	40S ribosomal protein S2 OS	0.39021111	0.21952736	0.0912853	0.04687208	FALSE	FALSE
O60506	SYNCRIP	Heterogeneous nuclear ribonucleopro	0.51329339	0.10531124	0.09171804	0.04685762	FALSE	TRUE
P55072	VCP	Transitional endoplasmic reticulum ATP	0.38386982	0.21748055	0.09240209	0.04697106	FALSE	TRUE
Q01105	SET	Protein SET OS	0.69722135	0.39718574	0.09328476	0.04718383	FALSE	TRUE
P35998	PSMC2	26S protease regulatory subunit 7 OS	0.467912	0.09994601	0.09543247	0.0475603	FALSE	FALSE
Q9H2J4	PDCL3	Phosducin-like protein 3 OS	0.86688367	0.18623745	0.09597619	0.04759795	FALSE	FALSE
P30050	RPL12	60S ribosomal protein L12 OS	0.70443438	0.40883236	0.09632708	0.04754006	FALSE	FALSE
P60842	EIF4A1	Eukaryotic initiation factor 4A-I OS	0.85086936	0.4973105	0.09750065	0.04788679	FALSE	FALSE
O15355	PPM1G	Protein phosphatase 1G OS	0.4869034	0.10634812	0.09755156	0.04768145	FALSE	FALSE
O75534	CSDE1	Cold shock domain-containing protein E	0.4095086	0.08981357	0.09794857	0.04764644	FALSE	TRUE
P16989	CSDA	DNA-binding protein A OS	0.8015366	0.47223168	0.09884481	0.04785344	FALSE	FALSE
P33992	MCM5	DNA replication licensing factor MCM5	0.61574019	0.13715809	0.09945704	0.04792164	FALSE	FALSE
Q9Y3F4	STRAP	Serine-threonine kinase receptor-assoc	0.42735396	0.09548391	0.09975454	0.04783826	FALSE	FALSE
Q58FF8	HSP90AB2P	Putative heat shock protein HSP 90-bet	0.81738756	0.4844621	0.099863	0.04766544	FALSE	FALSE
P50395	GDI2	Rab GDP dissociation inhibitor beta OS	0.4497802	0.10177456	0.10100397	0.04776157	FALSE	TRUE
P06753	TPM3	Tropomyosin alpha-3 chain OS	0.66274576	0.39928265	0.10268885	0.04833348	FALSE	FALSE
Q92841	DDX17	Probable ATP-dependent RNA helicase	0.71100407	0.16487021	0.10346366	0.04847376	FALSE	FALSE
P46776	RPL27A	60S ribosomal protein L27a OS	0.55982616	0.13004094	0.10364095	0.04833408	FALSE	FALSE
Q9UQE7	SMC3	Structural maintenance of chromosome	0.51423972	0.11998053	0.10409161	0.04832259	FALSE	FALSE
P33991	MCM4	DNA replication licensing factor MCM4	0.41271027	0.09644216	0.104251	0.0481766	FALSE	FALSE
Q8N766	KIAA0090	Uncharacterized protein KIAA0090 OS	0.45905283	0.10735956	0.10433506	0.04799728	FALSE	FALSE
Q14444	CAPRIN1	Caprin-1 OS	0.395091	0.09249096	0.10443518	0.04782692	FALSE	FALSE
Q9NY27	PPP4R2	Serine/threonine-protein phosphatase	0.47286342	0.28801189	0.10461221	0.04769316	FALSE	FALSE
Q9BQE3	TUBA1C	Tubulin alpha-1C chain OS	0.47790219	0.29204914	0.10520316	0.04774846	FALSE	FALSE
P13639	EEF2	Elongation factor 2 OS	0.40891191	0.25030785	0.10550243	0.04767147	FALSE	FALSE
Q9BVK6	TMED9	Transmembrane emp24 domain-contai	0.21460246	0.05076392	0.10550759	0.04746285	FALSE	TRUE
P10155	TROVE2	60 kDa SS-A/Ro ribonucleoprotein OS	0.20657568	0.04902814	0.10585297	0.04740845	FALSE	FALSE
Q86VP6	CAND1	Cullin-associated NEDD8-dissociated pr	0.41066848	0.09988202	0.10842594	0.04834782	FALSE	TRUE
P05141	SLC25A5	ADP/ATP translocase 2 OS	0.57071025	0.1391767	0.10870909	0.0482624	FALSE	FALSE
P23528	CFL1	Cofilin-1 OS	0.58234984	0.14300193	0.10944966	0.04837992	FALSE	FALSE
P55884	EIF3B	Eukaryotic translation initiation factor 3	0.40873871	0.10049299	0.10958119	0.04822837	FALSE	FALSE
P08865	RPSA	40S ribosomal protein SA OS	0.87864816	0.55047107	0.10971767	0.0480803	FALSE	FALSE
P67809	YBX1	Nuclease-sensitive element-binding pro	0.78793951	0.49399288	0.10984881	0.04793117	FALSE	FALSE
Q9Y262	EIF3L	Eukaryotic translation initiation factor 3	0.59313666	0.14772556	0.1109776	0.04821676	FALSE	FALSE
P27708	CAD	CAD protein OS	0.9741669	0.61486391	0.1110963	0.04806294	FALSE	FALSE
Q16531	DDB1	DNA damage-binding protein 1 OS	0.51150222	0.12795845	0.11145949	0.04801574	FALSE	FALSE
P06744	GPI	Glucose-6-phosphate isomerase OS	0.37706034	0.09465985	0.11184571	0.04777723	FALSE	FALSE
P46060	RANGAP1	Ran GTPase-activating protein 1 OS	0.64593606	0.16335189	0.1126506	0.04772005	FALSE	FALSE
Q13509	TUBB3	Tubulin beta-3 chain OS	0.34406867	0.08751232	0.11328464	0.04778951	FALSE	FALSE
Q15185	PTGES3	Prostaglandin E synthase 3 OS	0.74718531	0.19060736	0.11361383	0.04773033	FALSE	FALSE
P52701	MSH6	DNA mismatch repair protein Msh6 OS	0.53606463	0.13683935	0.1136862	0.04756419	FALSE	TRUE
P17812	CTPS	CTP synthase 1 OS	0.55729959	0.14240134	0.11379682	0.04741534	FALSE	FALSE
P62195	PSMC5	26S protease regulatory subunit 8 OS	0.46636066	0.12046787	0.11501462	0.04772716	FALSE	FALSE
Q9Y383	LUC7L2	Putative RNA-binding protein Luc7-like	0.73080536	0.18907795	0.1151935	0.04760707	FALSE	FALSE
P84103	SRSF3	Serine/arginine-rich splicing factor 3 OS	0.48770322	0.12667875	0.11563778	0.0475972	FALSE	FALSE
P63244	GNB2L1	Guanine nucleotide-binding protein sub	0.28301502	0.18306187	0.11574471	0.04744911	FALSE	FALSE
P23246	SFPQ	Splicing factor, proline- and glutamine-r	0.66981977	0.17636261	0.11718432	0.04784634	FALSE	FALSE
Q9NVI7	ATAD3A	ATPase family AAA domain-containing p	0.63534701	0.16860218	0.11808539	0.04802139	FALSE	FALSE

P31943	HNRNPH1	Heterogeneous nuclear ribonucleoprot	0.57312249	0.15253448	0.11842287	0.04796677	FALSE	FALSE
Q14974	KPNB1	Importin subunit beta-1 OS	0.44405385	0.29417976	0.12043244	0.04858716	FALSE	FALSE
P31153	MAT2A	S-adenosylmethionine synthase isoforn	0.44414365	0.12110968	0.1212616	0.04872831	FALSE	FALSE
P84098	RPL19	60S ribosomal protein L19 OS	0.51883668	0.14155649	0.12132804	0.04856306	FALSE	FALSE
P61978	HNRNPK	Heterogeneous nuclear ribonucleoprot	0.56337276	0.15537595	0.12261288	0.04888487	FALSE	FALSE
P06748	NPM1	Nucleophosmin OS	0.86292034	0.24102039	0.12413512	0.04929845	FALSE	FALSE
Q8WU90	ZC3H15	Zinc finger CCCH domain-containing pr	0.29026449	0.08118685	0.12430484	0.04917377	FALSE	FALSE
P30153	PPP2R1A	Serine/threonine-protein phosphatase	0.58767764	0.16477366	0.12460003	0.0490995	FALSE	FALSE
P61313	RPL15	60S ribosomal protein L15 OS	0.50086062	0.14223103	0.1261552	0.04952038	FALSE	FALSE
P39019	RPS19	40S ribosomal protein S19 OS	0.42178506	0.12186212	0.12829423	0.05016633	FALSE	FALSE
Q86UP2	KTN1	Kinectin OS	0.36426652	0.10526902	0.12832407	0.04998575	FALSE	TRUE
P11940	PABPC1	Polyadenylate-binding protein 1 OS	0.40390383	0.11776298	0.12943546	0.05022623	FALSE	FALSE
P06280	GLA	Alpha-galactosidase A OS	0.34837449	0.10262176	0.13073516	0.05053767	FALSE	TRUE
Q9Y5B9	SUPT16H	FACT complex subunit SPT16 OS	0.4652314	0.13734275	0.13101157	0.05045269	FALSE	FALSE
Q9Y678	COPG	Coatomer subunit gamma OS	0.366972	0.10857188	0.13128974	0.05036902	FALSE	TRUE
P12236	SLC25A6	ADP/ATP translocase 3 OS	0.56634707	0.16835818	0.1318984	0.0504123	FALSE	FALSE
Q00610	CLTC	Clathrin heavy chain 1 OS	0.47417165	0.14253574	0.13333294	0.05076972	FALSE	FALSE
P23588	EIF4B	Eukaryotic translation initiation factor 4	0.45172236	0.32012105	0.13445266	0.05081544	FALSE	FALSE
Q00059	TFAM	Transcription factor A, mitochondrial O	0.59730132	0.42440318	0.13502474	0.05084265	FALSE	FALSE
Q9Y295	DRG1	Developmentally-regulated GTP-bindin	0.35758247	0.10915898	0.13534292	0.0507744	FALSE	FALSE
O14818	PSMA7	Proteasome subunit alpha type-7 OS	0.34492211	0.10533824	0.13539788	0.05060828	FALSE	FALSE
O43707	ACTN4	Alpha-actinin-4 OS	0.60973849	0.1865386	0.13562795	0.05032424	FALSE	FALSE
P04083	ANXA1	Annexin A1 OS	0.48392382	0.34955261	0.13864776	0.05125766	FALSE	FALSE
P12004	PCNA	Proliferating cell nuclear antigen OS	0.46349594	0.1455713	0.13912436	0.0512475	FALSE	FALSE
P18615	RDBP	Negative elongation factor E OS	0.49612598	0.15676999	0.13994602	0.05136406	FALSE	FALSE
Q9NR30	DDX21	Nucleolar RNA helicase 2 OS	0.70093181	0.22167698	0.14006269	0.05122197	FALSE	FALSE
P83731	RPL24	60S ribosomal protein L24 OS	0.60519678	0.19178881	0.14033825	0.05113879	FALSE	FALSE
P39687	ANP32A	Acidic leucine-rich nuclear phosphopro	0.87320611	0.27747408	0.14070746	0.05109021	FALSE	TRUE
O60763	USO1	General vesicular transport factor p115	0.66902589	0.2132238	0.14111156	0.0510546	FALSE	TRUE
P62888	RPL30	60S ribosomal protein L30 OS	0.33530716	0.24537118	0.14156026	0.05103532	FALSE	FALSE
Q07021	C1QBP	Complement component 1 Q subcomp	0.21590847	0.15816988	0.14180658	0.05094347	FALSE	FALSE
P08758	ANXA5	Annexin A5 OS	0.33226512	0.24424799	0.14258515	0.05104281	FALSE	FALSE
P50914	RPL14	60S ribosomal protein L14 OS	0.15905709	0.11722073	0.1431639	0.05107016	FALSE	FALSE
Q9BTE3	MCMBP	Mini-chromosome maintenance compl	0.43232416	0.14161583	0.14490239	0.05150959	FALSE	FALSE
P13010	XRCC5	X-ray repair cross-complementing prot	0.25420415	0.08347443	0.1452471	0.05145223	FALSE	FALSE
P26358	DNMT1	DNA (cytosine-5)-methyltransferase 1 C	0.32588403	0.10950491	0.14851013	0.05242545	FALSE	FALSE
Q9BVA1	TUBB2B	Tubulin beta-2B chain OS	0.38316739	0.12969799	0.14956003	0.05261339	FALSE	FALSE
P10321	HLA-C	HLA class I histocompatibility antigen, C	0.49576827	0.16792691	0.14965857	0.05246651	FALSE	FALSE
Q01844	EWSR1	RNA-binding protein EWS OS	0.37904079	0.13108449	0.15268276	0.05316009	FALSE	FALSE
Q2NL82	TSR1	Pre-rRNA-processing protein TSR1 hom	0.6656885	0.23030442	0.15273887	0.05299813	FALSE	FALSE
Q9UQ80	PA2G4	Proliferation-associated protein 2G4 OS	0.39311656	0.30197359	0.15283817	0.0528522	FALSE	FALSE
Q08J23	NSUN2	tRNA (cytosine-5-)-methyltransferase N	0.68068466	0.2421806	0.15690539	0.05407474	FALSE	FALSE
P04406	GAPDH	Glyceraldehyde-3-phosphate dehydrog	0.4137447	0.32352155	0.15713497	0.05397091	FALSE	FALSE
Q9H3U1	UNC45A	Protein unc-45 homolog A OS	0.64448294	0.23043171	0.15764833	0.05396492	FALSE	FALSE
Q9UBQ5	EIF3K	Eukaryotic translation initiation factor 3	0.41028896	0.14736773	0.15834003	0.05401981	FALSE	FALSE
Q9BRL6	SRSF8	Serine/arginine-rich splicing factor 8 OS	0.4059115	0.14958379	0.16228051	0.05517899	FALSE	FALSE
Q9UMX5	NENF	Neudesin OS	0.59045059	0.21928787	0.16349283	0.0554059	FALSE	FALSE
P78371	CCT2	T-complex protein 1 subunit beta OS	0.52102825	0.41896438	0.16406853	0.05541628	FALSE	FALSE
Q8WUA2	PPIL4	Peptidyl-prolyl cis-trans isomerase-like	0.96255665	0.36025464	0.16470411	0.05544675	FALSE	FALSE
Q13310	PABPC4	Polyadenylate-binding protein 4 OS	0.50830918	0.19117895	0.16547756	0.05552327	FALSE	FALSE
P63173	RPL38	60S ribosomal protein L38 OS	0.69439372	0.26247827	0.16627135	0.0556061	FALSE	FALSE
Q15006	TTC35	Tetratricopeptide repeat protein 35 OS	0.29747855	0.11308834	0.1671785	0.05572617	FALSE	TRUE
P14868	DARS	Aspartyl-tRNA synthetase, cytoplasmic	0.46333875	0.17653012	0.1675309	0.05566114	FALSE	FALSE
Q14152	EIF3A	Eukaryotic translation initiation factor 3	0.56984367	0.2172144	0.16760922	0.05550577	FALSE	FALSE
P05204	HMGN2	Non-histone chromosomal protein HM	0.53549424	0.20472374	0.1680815	0.05548145	FALSE	FALSE
P00505	GOT2	Aspartate aminotransferase, mitochon	0.76935253	0.29615443	0.16918503	0.05548541	FALSE	FALSE
Q96DG6	CMBL	Carboxymethylenebutenolidase homol	0.57902947	0.22835735	0.17313453	0.0565981	FALSE	FALSE
Q15370	TCEB2	Transcription elongation factor B polyp	0.78564917	0.3107905	0.17363765	0.05658064	FALSE	FALSE
P19105	MYL12A	Myosin regulatory light chain 12A OS	0.63327865	0.25164023	0.17437914	0.05646034	FALSE	FALSE
P31689	DNAJA1	DnaJ homolog subfamily A member 1 O	0.329078	0.13120565	0.17494052	0.05646228	FALSE	FALSE
Q02878	RPL6	60S ribosomal protein L6 OS	0.4331334	0.17408139	0.17627621	0.05653443	FALSE	FALSE
Q02818	NUCB1	Nucleobindin-1 OS	0.47623561	0.4041174	0.17801699	0.05691319	FALSE	TRUE
P30041	PRDX6	Peroxiredoxin-6 OS	0.55261596	0.22575442	0.17902508	0.05705606	FALSE	FALSE
P62826	RAN	GTP-binding nuclear protein Ran OS	0.50779852	0.2078344	0.17934314	0.05697881	FALSE	FALSE
P33993	MCM7	DNA replication licensing factor MCM7	0.34583725	0.14169319	0.17951984	0.05685727	FALSE	FALSE
Q99714	HSD17B10	3-hydroxyacyl-CoA dehydrogenase type	0.67816503	0.28059649	0.18119962	0.05721106	FALSE	FALSE
P02768	ALB	Serum albumin OS	0.75575966	0.65315357	0.18294059	0.05758192	FALSE	TRUE
Q13409	DYNC1I2	Cytoplasmic dynein 1 intermediate chai	0.34244863	0.14660489	0.18713334	0.05835958	FALSE	FALSE
O75821	EIF3G	Eukaryotic translation initiation factor 3	0.5076309	0.22086186	0.19000603	0.05907425	FALSE	FALSE
P50991	CCT4	T-complex protein 1 subunit delta OS	0.45829238	0.40795807	0.19109001	0.05923013	FALSE	FALSE
P17844	DDX5	Probable ATP-dependent RNA helicase	0.70362965	0.31463424	0.19495983	0.06006338	FALSE	FALSE
P33176	KIF5B	Kinesin-1 heavy chain OS	0.48073322	0.21503663	0.19502156	0.05990088	FALSE	FALSE
P30048	PRDX3	Thioredoxin-dependent peroxide reduc	0.25196899	0.11341269	0.1961651	0.06007064	FALSE	FALSE
P46821	MAP1B	Microtubule-associated protein 1B OS	0.56888908	0.25621859	0.19627887	0.05992498	FALSE	FALSE
P49321	NASP	Nuclear autoantigenic sperm protein O	0.27294498	0.24765105	0.19647768	0.05980608	FALSE	FALSE
P07355	ANXA2	Annexin A2 OS	0.55900939	0.50863092	0.19727766	0.05987033	FALSE	FALSE
P60660	MYL6	Myosin light polypeptide 6 OS	0.90681969	0.41271057	0.19821204	0.05961999	FALSE	FALSE
P80303	NUCB2	Nucleobindin-2 OS	0.28957206	0.13405169	0.20139337	0.06039821	FALSE	TRUE
P58107	EPPK1	Epiplakin OS	0.64845592	0.30208883	0.20258328	0.06039873	FALSE	FALSE
P17987	TCP1	T-complex protein 1 subunit alpha OS	0.44520077	0.20829438	0.20339815	0.06046436	FALSE	TRUE
P18124	RPL7	60S ribosomal protein L7 OS	0.29944954	0.14089864	0.20447657	0.06043154	FALSE	FALSE
Q14683	SMC1A	Structural maintenance of chromosome	0.62126934	0.29384653	0.2054699	0.0605491	FALSE	FALSE
P31946	YWHAB	14-3-3 protein beta/alpha OS	0.63126191	0.30046604	0.2066834	0.06073067	FALSE	FALSE
P62081	RPS7	40S ribosomal protein S7 OS	0.45267761	0.21594409	0.20711209	0.06068126	FALSE	FALSE
Q99961	SH3GL1	Endophilin-A2 OS	0.41285414	0.19886363	0.20898673	0.06105455	FALSE	FALSE
O00571	DDX3X	ATP-dependent RNA helicase DDX3X O	0.58330907	0.28403153	0.21110252	0.06149596	FALSE	FALSE
Q99873	PRMT1	Protein arginine N-methyltransferase 1	0.79379354	0.38740582	0.21154998	0.06145023	FALSE	FALSE
P09622	DLD	Dihydrolipoyl dehydrogenase, mitocho	0.40828562	0.20308906	0.21531357	0.06236528	FALSE	FALSE
Q9UHV9	PFDN2	Prefoldin subunit 2 OS	0.36424061	0.18125651	0.21539746	0.06221234	FALSE	FALSE
Q96AG4	LRRC59	Leucine-rich repeat-containing protein	0.50342261	0.25143043	0.21612362	0.06224523	FALSE	TRUE
P13861	PRKAR2A	cAMP-dependent protein kinase type II	0.29755852	0.15097679	0.21929714	0.06280341	FALSE	FALSE
P54136	RARS	Arginyl-tRNA synthetase, cytoplasmic C	0.57128211	0.29046395	0.21971862	0.06274736	FALSE	FALSE
Q8TCJ2	STT3B	Dolichyl-diphosphooligosaccharide--pr	0.41256228	0.20992236	0.21987153	0.06261514	FALSE	TRUE
Q13347	EIF3I	Eukaryotic translation initiation factor 3	0.5170832	0.26373418	0.22035613	0.06257786	FALSE	FALSE

P26639	TARS	Threonyl-tRNA synthetase, cytoplasmic	0.65796662	0.33630062	0.22078586	0.06252524	FALSE	FALSE
P62829	RPL23	60S ribosomal protein L23 OS	0.66743901	0.34186287	0.22121568	0.06247295	FALSE	FALSE
P61289	PSME3	Proteasome activator complex subunit	0.29424313	0.15184107	0.22274193	0.06272972	FALSE	FALSE
P18206	VCL	Vinculin OS	0.526067	0.27152459	0.22278214	0.06256773	FALSE	FALSE
Q14696	MESDC2	LDLR chaperone MESD OS	0.65664261	0.34006031	0.22347173	0.0625885	FALSE	TRUE
P22234	PAICS	Multifunctional protein ADE2 OS	0.42052984	0.21915404	0.22476499	0.06277777	FALSE	FALSE
P24534	EEF1B2	Elongation factor 1-beta OS	0.38368522	0.2005497	0.22538124	0.06277742	FALSE	TRUE
Q9NQ39	RPS10P5	Putative 40S ribosomal protein S10-like	0.65732557	0.3528459	0.2309462	0.06397692	FALSE	FALSE
P06576	ATP5B	ATP synthase subunit beta, mitochondr	0.76255493	0.41257588	0.23261833	0.06426503	FALSE	FALSE
Q12905	ILF2	Interleukin enhancer-binding factor 2 C	0.51929226	0.28386917	0.23481528	0.06452132	FALSE	FALSE
O95347	SMC2	Structural maintenance of chromosome	0.56751814	0.31074859	0.23517193	0.06444514	FALSE	FALSE
P34932	HSPA4	Heat shock 70 kDa protein 4 OS	0.58844831	0.32300297	0.23569992	0.0644162	FALSE	FALSE
Q12931	TRAP1	Heat shock protein 75 kDa, mitochondr	0.41590241	0.22878324	0.23616241	0.06436956	FALSE	FALSE
O00159	MYO1C	Myosin-Ic OS	0.72756674	0.40311996	0.23771601	0.06461977	FALSE	FALSE
Q9H3P7	ACBD3	Golgi resident protein GCP60 OS	0.82460197	0.45898182	0.23870829	0.06471647	FALSE	TRUE
P32119	PRDX2	Peroxiredoxin-2 OS	0.37454457	0.39085224	0.23883004	0.06457727	FALSE	FALSE
Q5JRA6	MIA3	Melanoma inhibitory activity protein 3	0.80152898	0.44808745	0.23965517	0.06462849	FALSE	TRUE
P42704	LRPPRC	Leucine-rich PPR motif-containing prote	0.3061015	0.1731651	0.24224795	0.06515487	FALSE	FALSE
P26641	EEF1G	Elongation factor 1-gamma OS	0.17133182	0.18109605	0.24294954	0.06517116	FALSE	TRUE
O95816	BAG2	BAG family molecular chaperone regula	0.58766114	0.33355275	0.24297807	0.06500729	FALSE	FALSE
Q96QK1	VPS35	Vacuolar protein sorting-associated prc	0.86071801	0.48965734	0.24348206	0.06497115	FALSE	FALSE
P61981	YWHAG	14-3-3 protein gamma OS	0.65790541	0.37436426	0.24353259	0.06481452	FALSE	FALSE
P54886	ALDH18A1	Delta-1-pyrroline-5-carboxylate syntha	0.45552386	0.25995958	0.24417479	0.06481576	FALSE	FALSE
P15924	DSP	Desmoplakin OS	0.62793845	0.66692858	0.2445084	0.0647353	FALSE	FALSE
P54105	CLNS1A	Methylosome subunit pICln OS	0.87483865	0.50049799	0.24472451	0.06462422	FALSE	FALSE
P07814	EPRS	Bifunctional aminoacyl-tRNA synthetas	0.5910891	0.3403523	0.2461554	0.06483368	FALSE	FALSE
P29401	TKT	Transketolase OS	0.61645999	0.35764524	0.24783498	0.06510738	FALSE	FALSE
P67936	TPM4	Tropomyosin alpha-4 chain OS	0.71758599	0.41739693	0.24841599	0.06509182	FALSE	FALSE
O14654	IRS4	Insulin receptor substrate 4 OS	0.7989778	0.46566631	0.24886227	0.06504113	FALSE	FALSE
O76003	GLRX3	Glutaredoxin-3 OS	0.52821852	0.3089258	0.24963806	0.06507659	FALSE	FALSE
P62266	RPS23	40S ribosomal protein S23 OS	0.64878398	0.38488819	0.25286068	0.06574809	FALSE	FALSE
O43681	ASNA1	ATPase ASNA1 OS	0.8630124	0.51353505	0.2535509	0.06575938	FALSE	TRUE
Q9Y230	RUVBL2	RuvB-like 2 OS	0.69908457	0.41677846	0.25398219	0.06570362	FALSE	FALSE
P12956	XRCC6	X-ray repair cross-complementing prote	0.64360527	0.38392787	0.25411575	0.06557132	FALSE	FALSE
P61604	HSPE1	10 kDa heat shock protein, mitochondr	0.37172831	0.22175131	0.25412126	0.06540674	FALSE	FALSE
P47897	QARS	Glutaminyl-tRNA synthetase OS	0.6064911	0.36200643	0.25425321	0.06527544	FALSE	FALSE
P15311	EZR	Ezrin OS	0.77076266	0.46040968	0.25442756	0.06515567	FALSE	FALSE
P61106	RAB14	Ras-related protein Rab-14 OS	0.17297622	0.10399637	0.25590626	0.06536969	FALSE	TRUE
P45877	PPIC	Peptidyl-prolyl cis-trans isomerase C OS	0.78038395	0.47241853	0.25748644	0.06560849	FALSE	FALSE
P12268	IMPDH2	Inosine-5'-monophosphate dehy	0.57789937	0.35019499	0.25771939	0.06550368	FALSE	FALSE
O43175	PHGDH	D-3-phosphoglycerate dehydrogenase (	0.66445087	0.40370092	0.25832447	0.06549373	FALSE	FALSE
Q13283	G3BP1	Ras GTPase-activating protein-binding p	0.48085595	0.2948905	0.26048394	0.06587695	FALSE	FALSE
P59998	ARPC4	Actin-related protein 2/3 complex subu	0.76427476	0.47609766	0.26414213	0.06663635	FALSE	FALSE
P17655	CAPN2	Calpain-2 catalytic subunit OS	0.4242305	0.2656629	0.26537861	0.06678257	FALSE	TRUE
P37802	TAGLN2	Transgelin-2 OS	0.73532976	0.4646837	0.26752676	0.06715692	FALSE	FALSE
P12277	CKB	Creatine kinase B-type OS	0.62801008	0.39712048	0.26767987	0.06702985	FALSE	FALSE
P08574	CYC1	Cytochrome c1, heme protein, mitocho	0.61441732	0.39392075	0.27096671	0.0676862	FALSE	FALSE
P61163	ACTR1A	Alpha-centractin OS	0.52642297	0.33901499	0.27203697	0.06778699	FALSE	FALSE
P11586	MTHFD1	C-1-tetrahydrofolate synthase, cytoplas	0.6182979	0.40029562	0.27331024	0.06793775	FALSE	FALSE
Q14C86	GAPVD1	GTPase-activating protein and VPS9 do	0.55755997	0.3634138	0.27493757	0.06817558	FALSE	FALSE
P62633	CNBP	Cellular nucleic acid-binding protein OS	0.50809726	0.33560748	0.27816945	0.06864214	FALSE	TRUE
P53621	COPA	Coatomer subunit alpha OS	0.47712753	0.3159714	0.27880425	0.0686322	FALSE	TRUE
P25685	DNAJB1	DnaJ homolog subfamily B member 1 O	0.63983494	0.42899965	0.28184262	0.06921256	FALSE	FALSE
P18858	LIG1	DNA ligase 1 OS	0.43537442	0.29262163	0.28244146	0.06919249	FALSE	TRUE
P43487	RANBP1	Ran-specific GTPase-activating protein	0.67043968	0.45134267	0.28284127	0.06912387	FALSE	FALSE
P01605	1	Ig kappa chain V-I region Lay OS	1.48963474	1.00455549	0.28326672	0.06906183	FALSE	FALSE
Q9NR33	POLE4	DNA polymerase epsilon subunit 4 OS	0.34021597	0.23091839	0.28487009	0.06928658	FALSE	FALSE
P29692	EEF1D	Elongation factor 1-delta OS	0.52043278	0.35912237	0.28899347	0.06995477	FALSE	TRUE
P16949	STMN1	Stathmin OS	0.80407281	0.56002629	0.29133131	0.07035317	FALSE	FALSE
O00330	PDHX	Pyruvate dehydrogenase protein X com	0.0834479	0.05813112	0.29137795	0.07019769	FALSE	FALSE
O75369	FLNB	Filamin-B OS	0.65147619	0.46025777	0.29494306	0.0708886	FALSE	FALSE
P62280	RPS11	40S ribosomal protein S11 OS	0.8331374	0.58951762	0.29534033	0.07081667	FALSE	FALSE
P81605	DCD	Dermcidin OS	0.58622242	0.72476567	0.29623025	0.07086292	FALSE	FALSE
P35606	COPB2	Coatomer subunit beta' OS	0.6219476	0.44839701	0.30013524	0.07162852	FALSE	TRUE
Q96P70	IPO9	Importin-9 OS	0.44871115	0.32367168	0.30027104	0.07149311	FALSE	FALSE
Q96KC8	DNAJC1	DnaJ homolog subfamily C member 1 O	0.78164514	0.56526171	0.30092543	0.07131488	FALSE	TRUE
P48643	CCT5	T-complex protein 1 subunit epsilon OS	0.28101151	0.20466443	0.30275839	0.07158241	FALSE	FALSE
P06493	CDK1	Cyclin-dependent kinase 1 OS	0.68434988	0.49987631	0.30351446	0.07159467	FALSE	FALSE
P62857	RPS28	40S ribosomal protein S28 OS	0.87862303	0.64192203	0.30357145	0.07144235	FALSE	FALSE
Q5T749	KPRP	Keratinocyte proline-rich protein OS	0.51690414	0.65442568	0.3047144	0.07154572	FALSE	FALSE
Q9UHB9	SRP68	Signal recognition particle 68 kDa prote	0.55400316	0.40851302	0.30597721	0.07167669	FALSE	TRUE
Q9UBR2	CTSZ	Cathepsin Z OS	0.36385763	0.268784	0.30644515	0.07162128	FALSE	TRUE
Q8TEX9	IPO4	Importin-4 OS	0.57903245	0.43021952	0.30796073	0.07181042	FALSE	FALSE
Q15393	SF3B3	Splicing factor 3B subunit 3 OS	0.51007009	0.38198898	0.31003781	0.07212931	FALSE	FALSE
P24666	ACP1	Low molecular weight phosphotyrosine	0.31054108	0.2350379	0.31283351	0.07261356	FALSE	FALSE
P62820	RAB1A	Ras-related protein Rab-1A OS	0.40942564	0.31049238	0.31335648	0.07256927	FALSE	TRUE
P31151	S100A7	Protein S100-A7 OS	0.88552334	0.67639558	0.31526801	0.07284602	FALSE	TRUE
Q9P2J5	LARS	Leucyl-tRNA synthetase, cytoplasmic O	0.5104993	0.39269625	0.31714772	0.07311418	FALSE	FALSE
Q6IN85	SMEK1	Serine/threonine-protein phosphatase	0.84070239	0.65431852	0.32028662	0.07367075	FALSE	FALSE
P01034	CST3	Cystatin-C OS	0.65407623	0.51154212	0.32159238	0.07380412	FALSE	TRUE
P60510	PPP4C	Serine/threonine-protein phosphatase	0.78358598	0.61547911	0.32275686	0.07373846	FALSE	FALSE
Q16769	QPCT	Glutaminyl-peptide cyclotransferase OS	0.71333274	0.56121184	0.32319758	0.07367359	FALSE	FALSE
O60884	DNAJA2	DnaJ homolog subfamily A member 2 C	0.66806988	0.5256804	0.3232382	0.07351801	FALSE	FALSE
P42166	TMPO	Lamina-associated polypeptide 2, isofo	0.64820928	0.51083765	0.32365425	0.07344832	FALSE	FALSE
P61221	ABCE1	ATP-binding cassette sub-family E mem	0.86914878	0.69095199	0.32601954	0.07382031	FALSE	FALSE
P35232	PHB	Prohibitin OS	0.65552809	0.52307886	0.32703623	0.07388596	FALSE	FALSE
Q9Y285	FARSA	Phenylalanyl-tRNA synthetase alpha ch	0.51145143	0.41190959	0.32956529	0.07429225	FALSE	FALSE
Q9Y5S9	RBM8A	RNA-binding protein 8A OS	0.55439636	0.44806291	0.330525	0.07434375	FALSE	FALSE
Q9UBT2	UBA2	SUMO-activating enzyme subunit 2 OS	0.47766803	0.39002469	0.33334067	0.07481156	FALSE	FALSE
P48668	KRT6C	Keratin, type II cytoskeletal 6C OS	-0.1007302	0.08366087	0.33805563	0.07553624	FALSE	FALSE
O00410	IPO5	Importin-5 OS	0.80006131	0.67315704	0.34167104	0.07617666	FALSE	TRUE
P15586	GNS	N-acetylglucosamine-6-sulfatase OS	0.3418806	0.29082436	0.34474712	0.07652683	FALSE	FALSE
P30837	ALDH1B1	Aldehyde dehydrogenase X, mitochond	0.90275855	0.7837121	0.35049039	0.07763222	FALSE	FALSE
Q9NZZ3	CHMP5	Charged multivesicular body protein 5 (	0.45973898	0.40094763	0.35179323	0.0777514	FALSE	FALSE

Q9NQC3	RTN4	Reticulon-4 OS	0.94439655	0.82893443	0.35362272	0.07798621	FALSE	TRUE
P40926	MDH2	Malate dehydrogenase, mitochondrial (	0.83646419	0.73865445	0.3553512	0.07819778	FALSE	FALSE
O75396	SEC22B	Vesicle-trafficking protein SEC22b OS	0.62304186	0.5644728	0.36272283	0.07964756	FALSE	TRUE
Q96AE4	FUBP1	Far upstream element-binding protein :	0.40399437	0.37583623	0.37041994	0.08098787	FALSE	FALSE
P22314	UBA1	Ubiquitin-like modifier-activating enzyn	0.916668	0.86921586	0.3760218	0.08186056	FALSE	FALSE
Q92820	GGH	Gamma-glutamyl hydrolase OS	0.61737048	0.58547293	0.37605258	0.08169233	FALSE	FALSE
Q9BSJ8	ESYT1	Extended synaptotagmin-1 OS	0.72027244	0.69793031	0.38242101	0.08272227	FALSE	FALSE
P42167	TMPO	Lamina-associated polypeptide 2, isofo	0.64733392	0.62895382	0.3832246	0.0827201	FALSE	TRUE
P01617	1	Ig kappa chain V-II region TEW OS	1.03799457	1.0180187	0.386014	0.08314567	FALSE	FALSE
P62805	HIST1H4A	Histone H4 OS	0.33298705	0.53638417	0.39475475	0.08484862	FALSE	FALSE
P19012	KRT15	Keratin, type I cytoskeletal 15 OS	-0.3840047	0.38845082	0.39528788	0.08478397	FALSE	FALSE
P08559	PDHA1	Pyruvate dehydrogenase E1 componen	0.38113087	0.40555743	0.41065243	0.08789403	FALSE	FALSE
P11177	PDHB	Pyruvate dehydrogenase E1 componen	0.71886461	0.77083047	0.41300255	0.08821133	FALSE	FALSE
Q96KA5	CLPTM1L	Cleft lip and palate transmembrane pro	0.29015947	0.31129815	0.41316348	0.0880607	FALSE	FALSE
P12036	NEFH	Neurofilament heavy polypeptide OS	1.00305689	1.12631246	0.42721595	0.09086532	FALSE	FALSE
P13646	KRT13	Keratin, type I cytoskeletal 13 OS	0.57066788	1.0656316	0.45156197	0.09584301	FALSE	FALSE
Q8N1N4	KRT78	Keratin, type II cytoskeletal 78 OS	0.56568721	1.05836265	0.45229874	0.09579939	FALSE	FALSE
Q15435	PPP1R7	Protein phosphatase 1 regulatory subu	0.35304492	0.43189515	0.45401035	0.09596199	FALSE	FALSE
Q86Y46	KRT73	Keratin, type II cytoskeletal 73 OS	0.58242995	1.09967846	0.45579647	0.09613964	FALSE	FALSE
Q7Z3Y8	KRT27	Keratin, type I cytoskeletal 27 OS	1.12799099	1.3883911	0.45593891	0.09597058	FALSE	FALSE
Q9Y5M8	SRPRB	Signal recognition particle receptor sub	0.46277054	0.57076283	0.45658056	0.09570933	FALSE	TRUE
Q5D862	FLG2	Filaggrin-2 OS	0.53161282	1.00720414	0.45711919	0.09562507	FALSE	FALSE
P61019	RAB2A	Ras-related protein Rab-2A OS	0.41635088	0.51740772	0.45896597	0.09581425	FALSE	TRUE
P12273	PIP	Prolactin-inducible protein OS	1.1527707	1.43975286	0.4605449	0.09594685	FALSE	FALSE
P40925	MDH1	Malate dehydrogenase, cytoplasmic OS	0.64034057	0.82457578	0.47021642	0.09776142	FALSE	FALSE
P41091	EIF2S3	Eukaryotic translation initiation factor 2	0.17978953	0.24018095	0.48187636	0.09998115	FALSE	FALSE
P20042	EIF2S2	Eukaryotic translation initiation factor 2	0.1993072	0.4076507	0.48626264	0.10068575	FALSE	FALSE
P02538	KRT6A	Keratin, type II cytoskeletal 6A OS	0.46081475	0.97566393	0.49928109	0.10317123	FALSE	FALSE
P43307	SSR1	Translocon-associated protein subunit a	0.22162958	0.47022191	0.50005947	0.10312247	FALSE	TRUE
P35908	KRT2	Keratin, type II cytoskeletal 2 epiderma	0.53605006	1.14086487	0.50122787	0.10315418	FALSE	TRUE
Q04837	SSBP1	Single-stranded DNA-binding protein, n	0.22792356	0.48530566	0.50139806	0.10298075	FALSE	FALSE
O15371	EIF3D	Eukaryotic translation initiation factor 3	0.13210835	0.1973531	0.51743417	0.1060601	FALSE	FALSE
Q2M2I5	KRT24	Keratin, type I cytoskeletal 24 OS	0.91951548	1.3952735	0.52239855	0.10664763	FALSE	FALSE
P30086	PEBP1	Phosphatidylethanolamine-binding pro	0.63591459	0.96927343	0.52382197	0.10672392	FALSE	FALSE
P07339	CTSD	Cathepsin D OS	0.32393496	0.50777098	0.53270093	0.10831586	FALSE	FALSE
P13647	KRT5	Keratin, type II cytoskeletal 5 OS	0.45470063	1.06628034	0.53706022	0.10898428	FALSE	FALSE
O95678	KRT75	Keratin, type II cytoskeletal 75 OS	0.44560265	1.04949425	0.53863891	0.1090869	FALSE	FALSE
P05455	SSB	Lupus La protein OS	0.15057979	0.35586646	0.53988248	0.10912138	FALSE	FALSE
P04259	KRT6B	Keratin, type II cytoskeletal 6B OS	0.45263455	1.07130949	0.54042258	0.10901381	FALSE	FALSE
Q08188	TGM3	Protein-glutamine gamma-glutamyltrar	0.58121155	0.93803517	0.54192667	0.10910075	FALSE	FALSE
P13645	KRT10	Keratin, type I cytoskeletal 10 OS	0.53270165	1.2868408	0.54780122	0.11006546	FALSE	FALSE
Q00341	HDLBP	Vigilin OS	0.24113987	0.41831759	0.56457994	0.11321294	FALSE	FALSE
P12035	KRT3	Keratin, type II cytoskeletal 3 OS	0.41580922	1.06728989	0.56935868	0.11394645	FALSE	FALSE
P25311	AZGP1	Zinc-alpha-2-glycoprotein OS	0.83863508	1.4785743	0.56962052	0.11377489	FALSE	FALSE
Q96I24	FUBP3	Far upstream element-binding protein :	0.28174428	0.49780666	0.57028967	0.1136852	FALSE	FALSE
Q02413	DSG1	Desmoglein-1 OS	0.47884854	0.91360472	0.59392146	0.11816441	FALSE	FALSE
Q02543	RPL18A	60S ribosomal protein L18a OS	0.13026085	0.36747068	0.60176424	0.11949095	FALSE	FALSE
P05198	EIF2S1	Eukaryotic translation initiation factor 2	0.17834726	0.51796941	0.61143308	0.1211742	FALSE	FALSE
P01040	CSTA	Cystatin-A OS	0.66433402	1.3786548	0.61918658	0.12223424	FALSE	FALSE
Q14CN4	KRT72	Keratin, type II cytoskeletal 72 OS	0.38746138	1.18201135	0.62743413	0.12362236	FALSE	FALSE
O75533	SF3B1	Splicing factor 3B subunit 1 OS	-0.234346	0.54325551	0.6512727	0.12807103	FALSE	FALSE
Q04695	KRT17	Keratin, type I cytoskeletal 17 OS	0.37018575	1.25805289	0.66096011	0.12972512	FALSE	FALSE
P14923	JUP	Junction plakoglobin OS	0.48677162	1.17524688	0.66267142	0.12981039	FALSE	FALSE
P08779	KRT16	Keratin, type I cytoskeletal 16 OS	0.35320522	1.21289947	0.66407142	0.12983448	FALSE	FALSE
P0C0S8	HIST1H2AG	Histone H2A type 1 OS	0.11442735	0.27780319	0.66420588	0.12961151	FALSE	FALSE
P02533	KRT14	Keratin, type I cytoskeletal 14 OS	0.35803274	1.23817473	0.66616737	0.12974524	FALSE	FALSE
P29508	SERPINB3	Serpin B3 OS	0.39471516	1.0504961	0.68905292	0.13394592	FALSE	FALSE
Q7Z794	KRT77	Keratin, type II cytoskeletal 1b OS	0.2918461	1.180661	0.71024435	0.13753938	FALSE	FALSE
O00461	GOLIM4	Golgi integral membrane protein 4 OS	0.19820272	0.57493666	0.71121343	0.13746521	FALSE	TRUE
P05787	KRT8	Keratin, type II cytoskeletal 8 OS	0.27806527	1.13415246	0.71240941	0.13743509	FALSE	FALSE
Q8NC51	SERBP1	Plasminogen activator inhibitor 1 RNA-l	0.14301408	0.8521445	0.79866219	0.15378281	FALSE	FALSE
P05089	ARG1	Arginase-1 OS	0.11392656	0.49451054	0.79948746	0.15365071	FALSE	FALSE
P04264	KRT1	Keratin, type II cytoskeletal 1 OS	0.17133492	1.45645882	0.85739594	0.16446903	FALSE	FALSE
P46777	RPL5	60S ribosomal protein L5 OS	0.04302154	0.29839649	0.87195143	0.16694613	FALSE	FALSE
Q86YZ3	HRNR	Hornerin OS	-0.1139447	1.16956019	0.88151936	0.16846078	FALSE	FALSE
P82970	HMGN5	High mobility group nucleosome-bindir	0.04802358	0.64969181	0.90983867	0.17354646	FALSE	FALSE
Q9Y6E2	BZW2	Basic leucine zipper and W2 domain-co	0.01202817	0.57536511	0.98118411	0.18680472	FALSE	FALSE
P35527	KRT9	Keratin, type I cytoskeletal 9 OS	0.01496801	1.56237242	0.98826737	0.18780159	FALSE	FALSE