

Accession	Protein	Tissue location	MW [kDa]	pI	Scores	Peptides	SC	pat.1			pat.2			pat.3		
								iTRAQ (113/114)	Median (113/114)	CV[%] (113/114)	iTRAQ (115/116)	Median (115/116)	CV[%] (115/116)	iTRAQ (117/118)	Median (117/118)	CV[%] (117/118)
CH10_HUMAN	10 kDa heat shock protein, mitochondrial (Hsp10)	cellular	10.9	9.4	395	7	76.5	27	1.44	28.68	27	0.37	33.24	27	0.21	53.12
1433B_HUMAN	14-3-3 protein beta/alpha (Protein kinase C inhibitor)	cellular	28.1	4.6	280	7	26.8	20	1.46	15.66	20	0.45	21.86	20	0.46	33.22
1433Z_HUMAN	14-3-3 protein zeta/delta (Protein kinase C inhibitor)	cellular	27.7	4.6	260	9	24.9	16	1.49	18.68	16	0.44	21.17	16	0.49	29.58
PLCD1_HUMAN	1-phosphatidylinositol-4,5-bisphosphate phospholipase C	cellular	85.7	6.2	100	4	6.5	3	1.32	15.93	3	1.10	49.59	3	0.83	58.85
PRS6A_HUMAN	26S protease regulatory subunit 6A (Proteasome activator)	cellular	49.2	5.0	65	4	10.9	5	1.24	11.48	5	0.42	56.82	5	0.41	14.53
RS10_HUMAN	40S ribosomal protein S10 - Homo sapiens (Homo sapiens)	cellular	18.9	10.5	138	5	37.6	7	1.70	38.34	7	0.53	44.63	7	0.29	27.19
RS12_HUMAN	40S ribosomal protein S12 - Homo sapiens (Homo sapiens)	cellular	14.5	6.4	166	2	28.8	11	1.40	33.23	11	0.36	33.24	11	0.18	26.52
RS13_HUMAN	40S ribosomal protein S13 - Homo sapiens (Homo sapiens)	cellular	17.2	11.0	63	2	11.3	3	1.74	21.94	3	0.56	22.69	3	0.56	19.65
RS14_HUMAN	40S ribosomal protein S14 - Homo sapiens (Homo sapiens)	cellular	16.3	10.9	135	2	15.2	4	1.82	18.14	4	0.49	20.14	4	0.37	41.19
RS19_HUMAN	40S ribosomal protein S19 - Homo sapiens (Homo sapiens)	cellular	16.1	10.7	140	3	35.2	9	1.25	25.13	9	0.46	13.33	9	0.24	27.85
RS20_HUMAN	40S ribosomal protein S20 - Homo sapiens (Homo sapiens)	cellular	13.4	10.7	151	3	22.7	5	1.24	35.68	5	0.54	11.29	5	0.29	8.96
RS24_HUMAN	40S ribosomal protein S24 - Homo sapiens (Homo sapiens)	cellular	15.4	11.3	78	2	17.3	2	1.88	13.69	2	0.62	9.72	2	0.59	1.01
RS28_HUMAN	40S ribosomal protein S28 - Homo sapiens (Homo sapiens)	cellular	7.8	11.6	83	1	30.4	2	1.20	3.74	2	0.50	10.27	2	0.18	35.63
RS29_HUMAN	40S ribosomal protein S29 - Homo sapiens (Homo sapiens)	cellular	6.7	11.8	73	2	30.4	6	1.54	24.77	6	0.70	41.27	6	0.27	21.75
RS3A_HUMAN	40S ribosomal protein S3a - Homo sapiens (Homo sapiens)	cellular	29.9	10.4	97	2	10.2	3	1.65	14.76	3	0.56	5.22	3	0.41	34.19
RS6_HUMAN	40S ribosomal protein S6 (Phosphoprotein NP3)	cellular	28.7	11.5	116	3	12.4	4	1.24	15.69	4	0.57	31.70	4	0.47	18.60
RS8_HUMAN	40S ribosomal protein S8 - Homo sapiens (Homo sapiens)	cellular	24.2	11.0	130	4	17.3	5	1.63	24.15	5	0.41	30.79	5	0.43	40.35
RSSA_HUMAN	40S ribosomal protein SA (p40) (34/67 kDa lamina)	cellular	32.8	4.6	120	3	14.6	3	1.10	14.92	3	0.64	23.92	3	0.53	10.11
CH60_HUMAN	60 kDa heat shock protein, mitochondrial precursor	cellular	61.0	5.6	522	6	16.9	12	1.61	25.14	12	0.53	11.67	12	0.32	33.85
RLA1_HUMAN	60S acidic ribosomal protein P1 - Homo sapiens	cellular	11.5	4.0	95	1	14.0	3	1.68	10.31	3	0.38	16.73	3	0.19	10.95
RLA2_HUMAN	60S acidic ribosomal protein P2 (Renal carcinoma)	cellular	11.7	4.2	236	3	25.2	8	1.56	33.87	8	0.39	19.98	8	0.22	70.03
RL12_HUMAN	60S ribosomal protein L12 - Homo sapiens (Homo sapiens)	cellular	17.8	10.3	100	2	18.8	4	1.34	16.72	4	0.59	12.16	4	0.34	38.87
RL13_HUMAN	60S ribosomal protein L13 (Breast basic conserved)	cellular	24.2	12.2	69	2	8.5	2	1.27	38.65	2	0.48	3.70	2	0.34	23.18
RL18_HUMAN	60S ribosomal protein L18 - Homo sapiens (Homo sapiens)	cellular	21.6	12.3	84	3	25.0	3	1.56	32.92	3	0.34	93.95	3	0.51	33.90
RL3_HUMAN	60S ribosomal protein L3 (HIV-1 TAR RNA-binding)	cellular	46.1	10.8	45	1	2.2	2	1.21	12.09	2	0.43	41.02	2	0.52	12.85
RL36_HUMAN	60S ribosomal protein L36 - Homo sapiens (Homo sapiens)	cellular	12.2	12.2	60	2	21.0	2	1.86	49.42	2	0.56	2.13	2	0.50	70.56
RL4_HUMAN	60S ribosomal protein L4 (L1) - Homo sapiens (Homo sapiens)	cellular	47.7	11.8	102	3	11.2	3	1.40	11.62	3	0.61	39.21	3	0.49	31.96
RL5_HUMAN	60S ribosomal protein L5 - Homo sapiens (Homo sapiens)	cellular	34.3	10.2	119	5	13.8	5	1.35	26.94	5	0.53	13.61	5	0.38	27.05
RL6_HUMAN	60S ribosomal protein L6 (TAX-responsive enhancer)	cellular	32.7	11.1	56	2	6.6	3	1.99	31.64	3	0.35	31.87	2	0.51	9.38
RL7A_HUMAN	60S ribosomal protein L7a (Surfeit locus protein)	cellular	30.0	11.3	103	4	15.4	4	1.18	30.89	4	0.61	13.80	4	0.50	14.83
6PGD_HUMAN	6-phosphogluconate dehydrogenase, decarboxylase	cellular	53.1	7.0	102	2	7.2	2	1.47	4.09	2	0.81	31.12	2	0.42	17.33
GRP78_HUMAN	78 kDa glucose-regulated protein precursor (GRP78)	cellular	72.3	4.9	1288	25	44.2	54	1.37	31.69	54	0.45	36.70	54	0.25	51.78
AN32A_HUMAN	Acidic leucine-rich nuclear phosphoprotein 32 (AN32)	cellular	28.6	3.8	262	9	35.3	11	1.45	24.00	11	0.41	38.84	11	0.26	42.87
ACON_HUMAN	Aconitate hydratase, mitochondrial precursor (aconitase)	cellular	85.4	7.9	55	3	5.0	3	1.32	15.17	2	0.69	7.95	3	0.47	18.38
ACTC_HUMAN	Actin, alpha cardiac muscle 1 (Alpha-cardiac actin)	cellular	42.0	5.1	950	16	50.1	87	1.58	26.99	87	0.53	25.47	87	0.47	30.16
ACTB_HUMAN	Actin, cytoplasmic 1 (Beta-actin) - Homo sapiens	cellular	41.7	5.2	1616	21	58.1	111	1.60	30.93	111	0.52	25.27	109	0.45	37.83
ACTH_HUMAN	Actin, gamma-enteric smooth muscle (Smooth muscle)	cellular	41.8	5.2	944	16	50.3	79	1.55	27.04	79	0.53	25.62	79	0.47	32.12
ACBP_HUMAN	Acyl-CoA-binding protein (ACBP) (Diazepam-binding inhibitor)	cellular	10.0	6.2	411	6	74.7	16	1.48	23.89	16	0.52	31.41	16	0.24	48.18
CAP1_HUMAN	Adenyl cyclase-associated protein 1 (CAP 1) - Homo sapiens	cellular	51.8	9.1	89	2	3.4	3	1.85	4.71	3	0.47	58.96	3	0.41	56.91
CD38_HUMAN	ADP-ribosyl cyclase 1 (EC 3.2.2.5) (Cyclic ADP-ribose synthase)	cellular	34.3	9.2	30	1	4.3	2	1.68	52.13	2	1.02	29.09	2	0.30	2.54
AKA12_HUMAN	A-kinase anchor protein 12 (A-kinase anchor protein)	cellular	191.3	4.2	142	2	3.1	3	1.39	23.60	3	0.58	21.29	3	0.54	19.80
AK1A1_HUMAN	Alcohol dehydrogenase [NADP+] (EC 1.1.1.2) (Alcohol dehydrogenase)	cellular	36.5	6.4	75	2	9.5	2	1.57	13.34	2	0.90	14.70	2	0.55	19.30
ADH1B_HUMAN	Alcohol dehydrogenase 1B (EC 1.1.1.1) (Alcohol dehydrogenase)	cellular	39.8	9.7	70	2	8.5	2	1.74	26.57	2	2.29	101.74	2	1.23	18.72
ALDH2_HUMAN	Aldehyde dehydrogenase, mitochondrial precursor (aldehyde dehydrogenase)	cellular	56.3	6.8	115	4	7.5	7	1.20	38.12	7	1.90	37.76	7	0.79	55.79
CRYAB_HUMAN	Alpha crystallin B chain (Alpha(B)-crystallin) (Resilin)	cellular	20.1	6.9	151	3	16.0	8	1.23	13.51	8	1.72	23.12	8	1.29	34.01
A1AG1_HUMAN	Alpha-1-acid glycoprotein 1 precursor (AGP 1) (Haptoglobin)	blood	23.5	4.8	412	8	36.8	28	0.99	13.30	28	0.54	25.70	28	1.53	44.42
A1AG2_HUMAN	Alpha-1-acid glycoprotein 2 precursor (AGP 2) (Haptoglobin)	blood	23.6	4.9	348	8	32.3	15	1.04	19.98	15	0.49	23.85	15	1.59	53.58
AACT_HUMAN	Alpha-1-antichymotrypsin precursor (ACT) (Concussion)	blood	47.6	5.2	143	3	10.4	3	1.13	4.40	3	0.58	16.65	3	0.75	54.89
A1AT_HUMAN	Alpha-1-antitrypsin precursor (Alpha-1 protease inhibitor)	blood	46.7	5.3	1918	28	60.5	111	1.11	17.47	110	0.62	35.16	111	1.61	49.41
A1BG_HUMAN	Alpha-1B-glycoprotein precursor (Alpha-1-B glycoprotein)	blood	54.2	5.5	694	11	28.3	32	1.11	19.94	32	0.90	35.53	32	0.98	29.68
A2AP_HUMAN	Alpha-2-antiplasmin precursor (Alpha-2-plasmin inhibitor)	blood	54.5	5.9	70	2	7.5	4	1.14	28.52	4	0.69	22.45	3	0.78	17.44
FETUA_HUMAN	Alpha-2-HS-glycoprotein precursor (Fetuin-A) (Fetuin)	blood	39.3	5.4	852	12	39.0	30	1.03	18.04	30	0.75	23.38	30	1.27	52.06

A2MG_HUMAN	Alpha-2-macroglobulin precursor (Alpha-2-M)	blood	163.2	6.0	289	11	10.9	13	1.21	28.96	13	0.79	32.75	13	0.63	55.98
ACTN1_HUMAN	Alpha-actinin-1 (Alpha-actinin cytoskeletal isoform)	cellular	103.0	5.1	610	13	19.7	22	1.36	25.82	22	0.71	20.59	22	0.45	37.85
ACTN4_HUMAN	Alpha-actinin-4 (Non-muscle alpha-actinin 4) (F)	cellular	104.8	5.2	502	12	20.7	25	1.48	31.01	25	0.69	18.91	25	0.50	31.57
ENSA_HUMAN	Alpha-endosulfine (ARPP-19e) - Homo sapiens	cellular	13.4	7.5	55	2	24.0	2	1.15	10.20	2	0.57	22.78	2	0.28	34.72
ENOA_HUMAN	Alpha-enolase (EC 4.2.1.11) (2-phospho-D-glyce	cellular	47.1	7.7	995	18	47.5	40	1.76	30.40	40	0.71	25.89	40	0.21	32.92
AMBP_HUMAN	AMBP protein precursor [Contains: Alpha-1-mid	blood	39.0	5.9	219	4	14.5	12	1.20	17.92	12	0.83	23.32	12	0.72	27.22
ANXA1_HUMAN	Annexin A1 (Annexin I) (Lipocortin I) (Calpactin	cellular	38.7	6.7	931	15	49.7	33	1.50	27.00	33	0.83	27.29	33	0.54	27.30
ANXA2_HUMAN	Annexin A2 (Annexin II) (Lipocortin II) (Calpactin	extracellular	38.6	8.5	953	16	45.1	52	1.44	21.05	52	1.17	51.17	52	0.52	33.45
ANXA5_HUMAN	Annexin A5 (Annexin V) (Lipocortin V) (Endonex	cellular	35.9	4.8	470	9	31.6	24	1.38	22.60	24	0.97	26.16	23	0.46	15.67
ANXA6_HUMAN	Annexin A6 (Annexin VI) (Lipocortin VI) (P68) (P	cellular	75.8	5.3	259	7	12.5	10	1.53	26.18	10	1.04	23.79	9	0.60	20.27
AGR2_HUMAN	Anterior gradient protein 2 homolog precursor	extracellular	20.0	9.5	220	5	33.1	10	1.29	25.94	10	0.25	132.42	10	0.48	32.38
ANT3_HUMAN	Antithrombin-III precursor (ATIII) - Homo sapien	blood	52.6	6.3	255	6	17.7	12	1.28	19.73	12	0.99	36.35	12	0.93	34.41
APOA1_HUMAN	Apolipoprotein A-I precursor (Apo-AI) (ApoA-I)	blood	30.8	5.5	2074	34	78.3	168	1.03	15.68	168	0.80	30.13	168	1.43	44.95
APOA2_HUMAN	Apolipoprotein A-II precursor (Apo-AII) (ApoA-II)	blood	11.2	7.1	278	7	51.0	44	0.90	19.41	44	0.65	26.24	44	1.47	36.34
APOA4_HUMAN	Apolipoprotein A-IV precursor (Apo-AIV) (ApoA	blood	45.4	5.2	1288	25	56.1	67	1.09	17.15	67	0.66	22.88	67	0.92	39.46
APOC1_HUMAN	Apolipoprotein C-I precursor (Apo-CI) (ApoC-I)	blood	9.3	9.3	191	5	38.6	14	1.23	11.30	14	0.71	28.27	14	1.19	35.91
APOC3_HUMAN	Apolipoprotein C-III precursor (Apo-CIII) (ApoC	blood	10.8	5.1	133	1	16.2	5	0.89	12.69	5	1.07	17.70	5	2.01	18.75
APOD_HUMAN	Apolipoprotein D precursor (Apo-D) (ApoD) - H	blood	21.3	4.9	434	7	26.5	35	1.10	20.41	35	1.65	41.24	35	0.74	28.65
ARMET_HUMAN	ARMET protein precursor (Arginine-rich protein	extracellular	20.2	9.6	57	2	19.0	2	0.97	13.25	2	0.46	9.18	2	0.16	34.40
ASPN_HUMAN	Asporin precursor (Periodontal ligament-associat	extracellular	43.3	7.0	551	11	35.1	23	1.19	20.41	24	0.88	52.10	24	0.77	36.29
ATP5H_HUMAN	ATP synthase D chain, mitochondrial (EC 3.6.3.1)	cellular	18.5	5.1	41	2	11.8	3	1.96	51.01	3	0.56	34.43	3	0.36	10.50
ATPD_HUMAN	ATP synthase delta chain, mitochondrial precu	cellular	17.5	5.2	81	3	14.9	8	1.51	24.10	8	0.64	16.80	8	0.44	42.63
ATPA_HUMAN	ATP synthase subunit alpha, mitochondrial prec	cellular	59.7	9.6	479	8	22.6	23	1.21	19.43	23	0.78	22.09	23	0.52	30.45
ATPB_HUMAN	ATP synthase subunit beta, mitochondrial precu	cellular	56.5	5.1	352	10	26.3	17	1.40	16.38	17	0.91	42.30	17	0.56	35.21
ATIF1_HUMAN	ATPase inhibitor, mitochondrial precursor - Hort	cellular	12.2	9.9	44	2	18.9	2	1.39	0.38	2	0.40	21.92	2	0.25	22.97
B3AT_HUMAN	Band 3 anion transport protein (Anion exchange	cellular	101.7	4.9	116	3	6.0	4	1.26	17.45	4	1.29	44.98	4	0.71	37.78
BAF_HUMAN	Barrier-to-autointegration factor (Breakpoint cl	cellular	10.1	5.8	154	4	28.1	12	1.62	13.29	12	0.37	34.74	12	0.24	22.10
PGBM_HUMAN	Basement membrane-specific heparan sulfate	extracellular	468.5	6.1	287	9	3.2	10	1.47	28.36	10	1.16	33.10	10	0.65	39.18
APOH_HUMAN	Beta-2-glycoprotein 1 precursor (Beta-2-glycop	blood	38.3	9.5	309	6	26.1	14	1.37	14.97	14	0.53	28.52	14	0.95	29.34
BGLR_HUMAN	Beta-glucuronidase precursor (EC 3.2.1.31) (Bet	cellular	74.7	6.6	163	4	8.1	6	1.22	19.53	6	0.84	12.65	6	0.49	44.45
PGS1_HUMAN	Biglycan precursor (Bone/cartilage proteoglycan	extracellular	41.6	7.9	1308	16	45.4	89	1.14	14.66	88	0.54	34.10	89	0.97	25.26
KCD12_HUMAN	BTB/POZ domain-containing protein KCTD12 (P	cellular	35.7	5.4	55	2	7.7	3	1.28	8.05	3	1.00	14.88	3	0.58	34.17
CALD1_HUMAN	Caldesmon (CDM) - Homo sapiens (Human)	cellular	93.2	5.5	701	13	22.8	18	1.54	24.43	18	0.47	32.02	18	0.24	22.46
CALM_HUMAN	Calmodulin (CaM) - Homo sapiens (Human)	cellular	16.8	3.9	62	2	26.8	2	1.26	17.51	2	0.62	8.52	2	0.34	26.09
CALL5_HUMAN	Calmodulin-like protein 5 (Calmodulin-like skin	cellular	15.9	4.1	403	5	37.0	15	1.03	21.21	14	0.56	28.69	15	0.16	56.73
CALX_HUMAN	Calnexin precursor (Major histocompatibility co	cellular	67.5	4.3	417	8	17.1	17	1.65	20.58	17	0.47	50.97	17	0.26	38.15
CPNS1_HUMAN	Calpain small subunit 1 (CSS1) (Calcium-depend	cellular	28.3	4.9	46	2	6.0	3	1.29	10.66	3	0.69	17.97	3	0.47	27.09
CNN1_HUMAN	Calponin-1 (Calponin H1, smooth muscle) (Basid	cellular	33.1	9.7	136	4	17.5	7	1.72	16.26	7	0.59	26.34	7	0.50	15.24
CNN2_HUMAN	Calponin-2 (Calponin H2, smooth muscle) (Neut	cellular	33.7	7.7	127	5	14.6	6	1.40	24.69	6	0.60	33.18	6	0.34	33.93
CALR_HUMAN	Calreticulin precursor (CRP55) (Calregulin) (HAC	cellular	48.1	4.1	563	9	30.0	20	1.26	29.93	20	0.43	27.62	20	0.17	46.24
CALU_HUMAN	Calumenin precursor (Crocalbin) (IEF SSP 9302)	cellular	37.1	4.3	112	3	10.5	5	1.03	16.12	5	0.38	30.18	5	0.25	35.66
CAH2_HUMAN	Carbonic anhydrase 2 (EC 4.2.1.1) (Carbonic an	cellular	29.2	7.0	112	3	12.7	5	1.45	12.88	5	0.55	21.79	5	0.54	32.09
CBR1_HUMAN	Carbonyl reductase [NADPH] 1 (EC 1.1.1.184) (N	cellular	30.4	9.5	107	2	10.5	4	1.60	23.52	4	0.81	27.92	4	0.60	35.07
CATA_HUMAN	Catalase (EC 1.11.1.6) - Homo sapiens (Human)	cellular	59.7	7.0	236	5	16.3	6	1.07	18.61	6	1.23	33.40	6	0.73	27.69
COMT_HUMAN	Catechol O-methyltransferase (EC 2.1.1.6) - Hor	cellular	30.0	5.2	166	2	14.4	2	1.29	1.66	2	1.09	31.33	2	0.21	46.04
CTND1_HUMAN	Catenin delta-1 (p120 catenin) (p120(ctn)) (Cad	cellular	108.1	5.8	98	2	5.1	2	1.45	6.09	2	0.81	11.63	2	0.69	0.21
CATB_HUMAN	Cathepsin B precursor (EC 3.4.22.1) (Cathepsin	cellular	37.8	5.9	91	3	13.3	3	1.85	16.08	3	0.72	32.57	3	0.35	20.94
CATD_HUMAN	Cathepsin D precursor (EC 3.4.23.5) [Contains:	cellular	44.5	6.1	151	3	14.1	3	1.24	40.18	3	0.47	226.81	3	0.33	54.84
CAV1_HUMAN	Caveolin-1 - Homo sapiens (Human)	cellular	20.5	5.6	174	4	22.5	9	1.25	33.03	9	1.07	37.21	9	0.77	28.68
CD44_HUMAN	CD44 antigen precursor (Phagocytic glycoprotei	cellular	81.5	5.0	154	3	4.2	6	1.35	16.31	6	0.57	15.27	6	0.36	34.91
CD59_HUMAN	CD59 glycoprotein precursor (Membrane attack	cellular	14.2	6.1	89	2	11.7	2	1.81	10.89	2	1.34	8.93	2	0.70	45.04
CDC42_HUMAN	Cell division control protein 42 homolog precu	cellular	21.3	5.7	151	3	16.8	7	1.34	17.52	7	0.68	36.51	7	0.33	28.89
MUC18_HUMAN	Cell surface glycoprotein MUC18 precursor (Me	cellular	71.6	5.5	83	2	5.1	2	1.63	9.59	2	0.90	13.81	2	0.36	14.44
CERU_HUMAN	Ceruloplasmin precursor (EC 1.16.3.1) (Ferroxid	blood	122.1	5.4	122	3	5.7	5	1.10	21.05	5	1.04	23.82	5	0.66	26.37
CLIC1_HUMAN	Chloride intracellular channel protein 1 (Nuclea	cellular	26.9	4.9	55	2	6.2	9	1.61	21.63	9	0.43	6.35	9	0.42	31.08
CLIC4_HUMAN	Chloride intracellular channel protein 4 (Intrac	cellular	28.8	5.3	94	2	10.3	5	1.41	19.84	5	1.51	16.37	5	0.45	32.10
CBX3_HUMAN	Chromobox protein homolog 3 (Heterochromatin	cellular	20.8	5.1	46	2	11.5	2	1.54	23.06	2	0.55	42.66	2	1.04	173.93

CMA1_HUMAN	Chymase precursor (EC 3.4.21.39) (Mast cell protease)	extracellular	27.3	10.4	294	4	28.3	11	1.46	16.24	11	0.87	26.14	11	0.56	22.95
CLUS_HUMAN	Clusterin precursor (Complement-associated protein)	extracellular	52.5	5.9	457	11	26.7	24	1.32	27.77	24	0.93	34.36	24	0.62	41.36
RCL_HUMAN	c-Myc-responsive protein Rcl - Homo sapiens (H)	cellular	19.1	4.8	56	2	17.8	2	1.26	29.25	2	0.77	20.60	2	0.54	5.96
COF1_HUMAN	Cofilin-1 (Cofilin, non-muscle isoform) (18 kDa)	cellular	18.5	9.1	826	18	72.3	55	1.41	29.67	55	0.34	30.22	55	0.16	35.75
CO1A1_HUMAN	Collagen alpha-1(I) chain precursor (Alpha-1 type I)	extracellular	138.8	5.5	1663	22	25.0	149	0.89	21.01	149	0.43	51.26	148	0.55	30.51
CO3A1_HUMAN	Collagen alpha-1(III) chain precursor - Homo sapiens	extracellular	138.5	6.2	217	5	9.0	8	1.30	17.05	8	0.59	35.66	8	0.49	26.88
CO4A1_HUMAN	Collagen alpha-1(IV) chain precursor (Arrestin)	extracellular	160.5	9.4	78	2	2.3	2	1.61	11.16	2	2.47	68.90	2	0.71	53.96
CO5A1_HUMAN	Collagen alpha-1(V) chain precursor - Homo sapiens	extracellular	183.4	4.8	137	6	4.2	22	1.14	14.64	22	0.57	22.05	22	0.32	50.93
CO6A1_HUMAN	Collagen alpha-1(VI) chain precursor - Homo sapiens	extracellular	108.5	5.1	2119	31	32.2	152	0.88	19.36	152	1.10	37.55	149	0.73	34.58
COCA1_HUMAN	Collagen alpha-1(XII) chain precursor - Homo sapiens	extracellular	332.9	5.3	188	5	2.4	5	1.22	19.24	5	0.56	22.79	5	0.44	38.25
COEA1_HUMAN	Collagen alpha-1(XIV) chain precursor (Undulin)	extracellular	193.4	5.0	1942	34	22.6	98	1.14	19.62	98	1.05	38.60	94	0.62	32.88
COJA1_HUMAN	Collagen alpha-1(XIX) chain precursor (Collagenase)	extracellular	115.1	9.3	70	3	4.2	6	0.87	6.60	6	0.43	28.46	7	0.52	29.19
COFA1_HUMAN	Collagen alpha-1(XV) chain precursor [Contains: alpha-1(XV)]	extracellular	141.6	4.8	123	3	3.1	4	1.82	11.25	4	1.06	28.09	4	0.51	18.58
COIA1_HUMAN	Collagen alpha-1(XVIII) chain precursor [Contains: alpha-1(XVIII)]	extracellular	178.0	5.6	70	2	2.2	2	1.93	52.19	2	1.19	37.59	2	0.38	54.68
CO1A2_HUMAN	Collagen alpha-2(I) chain precursor (Alpha-2 type I)	extracellular	129.2	9.7	887	13	13.8	72	0.96	17.59	73	0.56	29.42	73	0.52	28.08
CO4A2_HUMAN	Collagen alpha-2(IV) chain precursor [Contains: alpha-2(IV)]	extracellular	167.4	9.6	257	3	6.1	9	1.30	20.91	9	1.14	35.82	9	0.64	42.39
CO5A2_HUMAN	Collagen alpha-2(V) chain precursor - Homo sapiens	extracellular	144.8	6.0	144	5	6.5	9	1.05	11.03	9	0.90	35.30	9	0.69	38.54
CO6A2_HUMAN	Collagen alpha-2(VI) chain precursor - Homo sapiens	extracellular	108.5	5.8	1545	25	28.0	101	0.79	19.88	101	1.01	34.80	101	0.70	41.63
CO6A3_HUMAN	Collagen alpha-3(VI) chain precursor - Homo sapiens	extracellular	343.3	6.4	7201	112	38.9	468	0.88	22.22	467	1.18	36.77	461	0.74	35.06
COMD8_HUMAN	COMM domain-containing protein 8 - Homo sapiens	cellular	21.1	5.2	44	1	6.6	2	1.54	44.62	2	0.60	5.61	2	0.28	25.98
C15_HUMAN	Complement C1s subcomponent precursor (EC 3.4.21.47)	blood	76.6	4.7	70	2	3.2	2	1.20	13.44	2	0.62	52.76	2	0.67	26.35
CO3_HUMAN	Complement C3 precursor [Contains: Complement C3]	blood	187.0	6.0	528	13	11.0	20	1.08	12.87	20	0.90	17.97	20	0.59	26.93
CO4B_HUMAN	Complement C4-B precursor (Basic complement component)	blood	192.7	6.8	211	6	5.7	7	1.19	22.22	7	1.19	29.32	7	0.54	59.61
CFAB_HUMAN	Complement factor B precursor (EC 3.4.21.47) (C3 convertase)	blood	85.5	6.7	165	3	6.9	4	0.89	18.45	4	0.90	24.85	4	0.97	28.78
CFAI_HUMAN	Complement factor I precursor (EC 3.4.21.45) (C3 inactivator)	blood	65.7	9.0	98	1	4.3	2	1.69	24.14	2	0.56	16.75	2	0.46	30.01
H2AY_HUMAN	Core histone macro-H2A.1 (Histone macroH2A.1)	cellular	39.6	10.3	198	2	10.5	3	1.48	24.15	3	0.59	18.58	3	0.66	16.25
CLC11_HUMAN	C-type lectin domain family 11 member A precursor	cellular	35.7	4.9	44	2	7.7	2	1.12	3.65	2	0.81	48.67	2	0.56	18.46
CDN2B_HUMAN	Cyclin-dependent kinase 4 inhibitor B (p14-INK4)	extracellular	14.7	6.1	40	2	17.4	9	0.70	19.83	9	0.88	26.71	9	0.73	31.30
CYTB_HUMAN	Cystatin-B (Stefin-B) (Liver thiol proteinase inhibitor)	cellular	11.1	7.9	130	3	30.6	6	1.47	11.94	6	0.68	69.73	6	0.40	33.42
CSR1P1_HUMAN	Cysteine and glycine-rich protein 1 (Cysteine-rich protein)	cellular	20.6	10.0	357	3	40.4	6	1.60	18.37	6	0.86	36.70	6	0.31	88.14
CRIP1_HUMAN	Cysteine-rich protein 1 (Cysteine-rich intestinal protein)	cellular	8.5	10.2	58	2	27.3	6	1.58	21.68	6	0.66	17.31	6	0.15	49.12
CRIP2_HUMAN	Cysteine-rich protein 2 (CRP2) (Protein ESP1) - Homo sapiens	cellular	22.5	10.1	186	5	34.6	7	1.96	22.37	7	0.57	19.30	7	0.32	27.14
CYB5_HUMAN	Cytochrome b5 - Homo sapiens (Human)	cellular	15.3	4.7	206	2	35.8	8	1.47	22.15	8	0.85	24.17	8	0.39	23.91
CYB5B_HUMAN	Cytochrome b5 type B precursor (Cytochrome b5)	cellular	16.3	4.7	62	1	6.2	2	1.03	47.91	2	0.76	38.47	2	0.34	51.14
CYC_HUMAN	Cytochrome c - Homo sapiens (Human)	cellular	11.7	10.1	102	3	24.8	7	1.55	9.93	7	0.57	35.30	7	0.25	34.56
COX5B_HUMAN	Cytochrome c oxidase subunit 5B, mitochondrial	cellular	13.7	10.2	72	3	31.0	3	1.42	6.20	3	0.72	22.01	3	0.57	14.04
CX6B1_HUMAN	Cytochrome c oxidase subunit VIb isoform 1 (EC 1.10.3.2)	cellular	10.2	7.5	92	2	26.7	3	1.46	7.34	3	0.42	21.19	3	0.38	11.78
CKAP4_HUMAN	Cytoskeleton-associated protein 4 (63 kDa member)	cellular	66.0	5.6	239	6	14.0	7	1.38	24.44	7	0.45	40.44	7	0.36	46.27
AMPL_HUMAN	Cytosol aminopeptidase (EC 3.4.11.1) (Leucine aminopeptidase)	cellular	56.1	9.0	87	2	6.4	2	1.35	5.87	2	0.78	31.88	2	0.34	17.35
CNDP2_HUMAN	Cytosolic nonspecific dipeptidase (CNDP dipeptidase)	cellular	52.8	5.6	143	4	10.9	4	1.63	10.96	4	0.57	14.45	4	0.50	31.28
CYTA_HUMAN	Cytospin-A (SPECC1-like protein) (Renal carcinoma)	cellular	124.5	5.5	30	1	0.6	7	1.11	8.50	7	0.68	15.97	7	0.88	15.56
SERA_HUMAN	D-3-phosphoglycerate dehydrogenase (EC 1.1.1.41)	cellular	56.6	6.3	50	2	3.6	2	1.32	5.59	2	0.79	28.82	2	0.62	29.87
DOPD_HUMAN	D-dopachrome decarboxylase (EC 4.1.1.84) (D-dopachrome decarboxylase)	cellular	12.7	7.8	110	3	27.1	3	1.07	35.15	3	0.64	18.74	3	0.70	13.54
PGS2_HUMAN	Decorin precursor (Bone proteoglycan II) (PG-S2)	extracellular	39.7	9.5	1368	24	60.2	141	1.10	35.77	141	0.99	27.01	139	0.99	39.43
DERM_HUMAN	Dermatopontin precursor (Tyrosine-rich acidic glycoprotein)	extracellular	24.0	4.5	180	2	13.9	4	0.83	18.99	4	1.46	7.57	4	1.59	54.11
DEST_HUMAN	Destrin (Actin-depolymerizing factor) (ADF) - Homo sapiens	cellular	18.5	9.2	99	3	17.0	4	1.58	13.84	4	0.50	12.65	4	0.29	25.68
DPYL3_HUMAN	Dihydropyrimidinase-related protein 3 (DRP-3)	cellular	61.9	6.0	144	3	6.0	5	1.27	28.91	5	1.00	12.40	5	0.52	36.92
APEX1_HUMAN	DNA-(apurinic or apyrimidinic site) lyase (EC 4.2.1.10)	cellular	35.5	9.2	71	2	6.6	4	1.22	19.14	4	0.81	16.52	4	0.39	17.55
DJB11_HUMAN	DnaJ homolog subfamily B member 11 precursor	cellular	40.5	5.8	47	2	5.9	2	1.54	2.71	2	0.67	16.48	2	0.44	58.55
DCTN2_HUMAN	Dynactin subunit 2 (Dynactin complex 50 kDa subunit)	cellular	44.2	5.0	110	3	11.2	5	1.50	12.25	5	0.66	16.91	5	0.43	18.97
EHD2_HUMAN	EH domain-containing protein 2 - Homo sapiens	cellular	61.1	6.0	281	7	22.3	11	1.49	30.14	11	1.21	32.50	11	0.73	34.43
EF1A1_HUMAN	Elongation factor 1-alpha 1 (EF-1-alpha-1) (Elongation factor)	cellular	50.1	9.7	622	10	22.3	20	1.50	28.75	20	0.60	28.60	20	0.31	45.99
EF1B_HUMAN	Elongation factor 1-beta (EF-1-beta) - Homo sapiens	cellular	24.7	4.3	194	5	28.9	5	0.99	49.40	5	0.33	32.11	4	0.20	35.99
EF1D_HUMAN	Elongation factor 1-delta (EF-1-delta) (Antigen H)	cellular	31.1	4.8	317	7	25.3	9	1.62	25.94	9	0.46	42.48	9	0.17	34.80
EF2_HUMAN	Elongation factor 2 (EF-2) - Homo sapiens (Human)	cellular	95.3	6.4	184	5	8.2	8	1.05	20.63	8	0.93	31.82	8	0.43	40.01
EFTU_HUMAN	Elongation factor Tu, mitochondrial precursor (EFTu)	cellular	49.5	7.9	188	4	13.5	4	1.66	29.85	4	0.75	25.05	4	0.42	43.27
EMIL1_HUMAN	EMILIN-1 precursor (Elastin microfibril interface-associated protein)	extracellular	106.6	5.0	415	11	16.0	13	1.57	20.06	13	0.84	24.03	13	0.51	29.12

ERP29_HUMAN	Endoplasmic reticulum protein Erp29 precursor	cellular	29.0	7.5	158	5	22.6	7	1.33	27.41	7	0.43	42.80	7	0.31	45.01
ENPL_HUMAN	Endoplasmic precursor (Heat shock protein 90)	cellular	92.4	4.6	412	11	15.1	29	1.39	19.64	29	0.50	35.25	29	0.33	27.72
ECHM_HUMAN	Enoyl-CoA hydratase, mitochondrial precursor	cellular	31.4	9.4	168	2	10.3	9	1.36	17.91	9	0.84	28.38	9	0.41	28.13
IF4H_HUMAN	Eukaryotic translation initiation factor 4H (eIF-4)	cellular	27.4	7.5	41	2	7.7	2	1.75	11.05	2	0.62	28.12	2	0.10	35.53
IF5A1_HUMAN	Eukaryotic translation initiation factor 5A-1 (eIF)	cellular	16.8	4.9	285	4	33.8	10	1.63	35.10	11	0.44	37.91	11	0.22	27.26
ECM1_HUMAN	Extracellular matrix protein 1 precursor (Secret)	extracellular	60.6	6.3	47	2	4.8	2	0.73	0.10	2	0.84	13.62	2	0.56	4.87
SODE_HUMAN	Extracellular superoxide dismutase [Cu-Zn] pre	extracellular	25.8	6.2	168	4	22.1	12	1.06	25.36	12	1.28	35.60	12	0.80	56.81
NHERF_HUMAN	Ezrin-radixin-moesin-binding phosphoprotein 5	cellular	38.8	5.5	160	4	20.9	5	1.53	12.56	5	0.46	48.22	5	0.18	33.65
FUBP1_HUMAN	Far upstream element-binding protein 1 (FUSE)	cellular	67.5	7.8	274	9	18.9	13	1.20	33.85	13	0.61	37.81	13	0.40	24.86
FAS_HUMAN	Fatty acid synthase (EC 2.3.1.85) [Includes: [Ac]	cellular	273.2	6.0	61	2	1.8	3	1.22	2.21	3	0.82	42.88	3	0.43	13.67
FABPA_HUMAN	Fatty acid-binding protein, adipocyte (AFABP)	cellular	14.7	7.5	326	6	47.7	18	1.38	28.27	18	1.89	26.92	18	1.24	34.15
FABPE_HUMAN	Fatty acid-binding protein, epidermal (E-FABP)	cellular	15.2	7.5	100	2	17.0	12	1.22	21.93	12	0.84	17.93	12	0.54	15.48
FRIH_HUMAN	Ferritin heavy chain (EC 1.16.3.1) (Ferritin H sub	cellular	21.2	5.2	143	4	22.4	7	1.55	13.86	7	0.69	16.39	7	0.39	18.95
FRIL_HUMAN	Ferritin light chain (Ferritin L subunit) - Homo sa	cellular	20.0	5.4	278	4	26.3	11	1.59	22.88	11	0.73	14.01	11	0.38	24.58
FBN1_HUMAN	Fibrillin-1 precursor - Homo sapiens (Human)	extracellular	312.1	4.7	1006	26	10.7	56	1.29	29.94	56	1.39	34.49	56	0.39	40.57
FIBA_HUMAN	Fibrinogen alpha chain precursor [Contains: Fibr	blood	94.9	5.6	1258	25	35.5	65	1.46	23.64	65	0.88	31.64	65	0.86	32.14
FIBB_HUMAN	Fibrinogen beta chain precursor [Contains: Fibr	blood	55.9	9.3	775	16	41.8	40	1.47	23.30	40	0.89	28.50	40	0.82	38.99
FIBG_HUMAN	Fibrinogen gamma chain precursor - Homo sapi	blood	51.5	5.3	506	13	24.9	20	1.27	29.56	20	1.14	37.86	20	1.26	56.95
FMOD_HUMAN	Fibromodulin precursor (FM) (Collagen-binding	extracellular	43.2	5.5	88	3	6.4	3	1.41	8.26	3	0.85	7.24	3	0.48	12.46
FINC_HUMAN	Fibronectin precursor (FN) (Cold-insoluble glob	extracellular	262.4	5.4	588	17	11.7	28	1.17	28.16	28	0.69	40.52	28	0.36	41.40
FBLN1_HUMAN	Fibulin-1 precursor - Homo sapiens (Human)	extracellular	77.2	5.0	103	4	8.5	4	1.26	19.39	4	1.01	46.19	4	0.59	50.72
FLNA_HUMAN	Filamin-A (Alpha-filamin) (Filamin-1) (Endotheli	cellular	280.6	5.7	3247	56	31.9	112	1.50	24.18	112	0.64	26.85	111	0.33	37.09
FLNB_HUMAN	Filamin-B (FLN-B) (Beta-filamin) (Actin-binding	cellular	278.0	5.4	481	12	8.1	17	1.44	23.43	17	0.63	29.02	16	0.39	37.35
FKB1A_HUMAN	FK506-binding protein 1A (EC 5.2.1.8) (Peptidyl-	cellular	11.9	9.1	201	2	34.3	6	1.35	24.73	6	0.47	16.85	6	0.45	39.99
FKBP2_HUMAN	FK506-binding protein 2 precursor (EC 5.2.1.8)	cellular	15.6	10.1	52	2	14.1	3	1.51	42.87	3	0.59	27.80	3	0.31	12.57
FKBP3_HUMAN	FK506-binding protein 3 (EC 5.2.1.8) (Peptidyl-p	cellular	25.2	9.8	31	1	3.6	2	1.11	17.29	2	0.56	9.47	2	0.21	0.32
BLVRB_HUMAN	Flavin reductase (EC 1.5.1.30) (FR) (NADPH-dep	cellular	22.1	7.9	154	3	23.3	6	1.48	25.28	6	0.84	30.78	6	0.43	10.13
FHL1_HUMAN	Four and a half LIM domains protein 1 (FHL-1)	cellular	36.2	10.5	193	6	15.2	6	1.36	22.96	6	1.35	25.25	6	0.98	7.44
ALDOA_HUMAN	Fructose-bisphosphate aldolase A (EC 4.1.2.13)	cellular	39.4	9.2	663	10	39.3	25	1.55	24.87	25	0.54	21.32	25	0.28	26.03
ALDOC_HUMAN	Fructose-bisphosphate aldolase C (EC 4.1.2.13)	cellular	39.4	6.5	163	2	11.0	4	2.53	26.69	4	0.77	30.19	4	0.26	32.80
FUSIP_HUMAN	FUS-interacting serine-arginine-rich protein 1	cellular	31.3	11.8	92	3	18.7	3	1.27	35.50	3	0.54	1.13	3	0.23	11.62
LEG1_HUMAN	Galectin-1 (Lectin galactoside-binding soluble 1)	cellular	14.7	5.2	725	10	63.7	42	1.49	31.78	42	0.50	27.33	42	0.37	25.09
LEG3_HUMAN	Galectin-3 (Galactose-specific lectin 3) (Mac-2	cellular	26.2	9.1	384	6	30.8	20	1.40	26.29	20	0.42	18.24	20	0.25	37.74
GGT5_HUMAN	Gamma-glutamyltransferase 5 precursor (EC 2.3.	cellular	62.2	8.0	52	2	4.3	7	1.45	12.77	7	0.63	12.00	6	0.23	10.24
SYUG_HUMAN	Gamma-synuclein (Persyn) (Breast cancer-spec	cellular	13.3	4.7	60	2	19.7	2	1.66	12.61	2	1.13	3.25	2	0.87	0.84
GELS_HUMAN	Gelsolin precursor (Actin-depolymerizing factor	cellular	85.6	5.9	676	15	23.8	38	1.21	22.47	38	0.70	28.55	38	0.76	28.62
G6PD_HUMAN	Glucose-6-phosphate 1-dehydrogenase (EC 1.1.	cellular	59.2	6.4	64	2	4.7	3	1.26	3.19	3	1.22	31.38	3	0.78	39.47
G6PI_HUMAN	Glucose-6-phosphate isomerase (EC 5.3.1.9) (G	cellular	63.1	9.1	91	1	3.0	2	1.54	17.23	2	1.69	24.40	2	0.41	11.66
GLU2B_HUMAN	Glucosidase 2 subunit beta precursor (Glucosid	cellular	59.4	4.2	234	8	13.3	12	1.68	36.49	12	0.46	28.54	12	0.25	51.16
GLRX1_HUMAN	Glutaredoxin-1 (Thioltransferase-1) (TTase-1) -	cellular	11.8	9.7	102	2	32.1	5	1.29	25.06	5	0.41	38.34	5	0.23	15.40
GSTP1_HUMAN	Glutathione S-transferase P (EC 2.5.1.18) (GST	cellular	23.3	5.3	346	3	34.8	8	2.28	39.25	8	0.88	14.48	8	0.57	26.72
GSTO1_HUMAN	Glutathione transferase omega-1 (EC 2.5.1.18)	cellular	27.5	6.3	161	5	19.9	5	1.93	27.37	5	0.52	24.14	5	0.39	17.09
G3P_HUMAN	Glyceraldehyde-3-phosphate dehydrogenase (E	cellular	36.0	9.3	671	11	38.2	31	1.60	22.82	31	0.89	20.64	31	0.24	40.85
GPDA_HUMAN	Glycerol-3-phosphate dehydrogenase [NAD+], c	cellular	37.5	5.8	310	6	26.4	12	1.27	28.86	12	1.56	27.38	12	0.88	29.56
HPT_HUMAN	Haptoglobin precursor [Contains: Haptoglobin a	blood	45.2	6.1	910	15	40.6	41	1.15	20.15	41	0.87	24.95	41	1.38	35.85
HSP71_HUMAN	Heat shock 70 kDa protein 1 (HSP70.1) (HSP70-	cellular	70.0	5.4	1431	21	38.2	70	1.54	27.35	70	0.58	27.32	69	0.45	36.01
HSP7C_HUMAN	Heat shock cognate 71 kDa protein (Heat shock	cellular	70.9	5.2	1075	15	33.6	40	1.50	25.17	40	0.54	30.67	40	0.36	30.11
HS90A_HUMAN	Heat shock protein HSP 90-alpha (HSP 86) (Ren	cellular	84.6	4.8	336	8	14.6	15	1.49	30.15	15	0.58	47.45	15	0.33	34.62
HS90B_HUMAN	Heat shock protein HSP 90-beta (HSP 84) (HSP	cellular	83.2	4.8	418	11	17.4	21	1.35	21.94	21	0.62	42.31	21	0.40	35.28
HSPB1_HUMAN	Heat-shock protein beta-1 (HspB1) (Heat shock	cellular	22.8	6.0	505	9	52.2	33	1.04	32.97	34	1.01	35.81	34	0.36	32.94
HSPB6_HUMAN	Heat-shock protein beta-6 (HspB6) (Heat-shock	cellular	17.1	6.0	163	4	26.2	6	1.28	15.95	6	0.98	23.92	6	0.65	20.78
HCLS1_HUMAN	Hematopoietic lineage cell-specific protein (Her	cellular	54.0	4.6	55	2	4.1	3	1.16	5.81	3	0.66	4.07	3	0.36	27.72
HEBP2_HUMAN	Heme-binding protein 2 (Protein SOUL) (Placen	cellular	22.9	4.4	61	1	6.3	2	1.45	44.64	2	0.63	6.97	2	0.25	5.08
HBA_HUMAN	Hemoglobin subunit alpha (Hemoglobin alpha ch	blood	15.2	9.4	623	11	74.6	88	1.58	30.42	87	1.75	40.03	89	1.12	29.81
HBB_HUMAN	Hemoglobin subunit beta (Hemoglobin beta cha	blood	16.0	6.9	1343	16	88.4	127	1.53	27.90	129	1.49	60.30	127	1.09	37.60
HBD_HUMAN	Hemoglobin subunit delta (Hemoglobin delta ch	blood	16.0	9.1	1110	14	91.2	77	1.46	30.78	77	1.71	42.11	76	1.11	39.40
HEMO_HUMAN	Hemopexin precursor (Beta-1B-glycoprotein) -	blood	51.6	6.6	918	17	30.5	68	1.00	27.85	68	0.91	23.88	68	1.49	50.26

HDGF_HUMAN	Hepatoma-derived growth factor (HDGF) (High-mobility group protein B1)	cellular	26.8	4.6	105	4	19.2	5	1.87	17.13	5	0.40	22.90	5	0.22	50.74
ROAA_HUMAN	Heterogeneous nuclear ribonucleoprotein A/B	cellular	36.6	9.5	115	4	10.3	8	1.37	23.13	8	0.43	17.01	8	0.32	36.25
ROA1_HUMAN	Heterogeneous nuclear ribonucleoprotein A1	cellular	38.8	9.7	674	13	28.8	29	1.40	21.23	29	0.39	47.44	29	0.28	34.66
ROA3_HUMAN	Heterogeneous nuclear ribonucleoprotein A3	cellular	39.6	9.6	554	13	31.2	26	1.33	17.54	26	0.42	21.81	26	0.26	36.89
HNRPD_HUMAN	Heterogeneous nuclear ribonucleoprotein D0	cellular	38.4	8.5	434	8	27.0	21	1.38	18.09	21	0.47	29.73	21	0.33	41.46
HNRH1_HUMAN	Heterogeneous nuclear ribonucleoprotein H	cellular	49.2	5.9	269	7	19.4	15	1.77	24.48	15	0.58	47.69	15	0.40	29.83
HNRH3_HUMAN	Heterogeneous nuclear ribonucleoprotein H3	cellular	36.9	6.4	159	2	7.8	3	1.38	8.85	3	0.84	10.36	3	0.26	10.09
HNRPK_HUMAN	Heterogeneous nuclear ribonucleoprotein K	cellular	50.9	5.3	332	7	19.4	15	1.52	17.59	15	0.45	28.55	15	0.33	44.21
HNRPM_HUMAN	Heterogeneous nuclear ribonucleoprotein M	cellular	77.5	9.6	236	5	15.8	8	1.68	18.67	8	0.56	28.92	8	0.34	26.23
HNRPU_HUMAN	Heterogeneous nuclear ribonucleoprotein U	cellular	90.5	5.7	253	6	9.7	11	1.58	21.80	11	0.67	36.36	11	0.43	29.58
ROA2_HUMAN	Heterogeneous nuclear ribonucleoproteins A2/B	cellular	37.4	9.3	1073	21	52.7	62	1.55	25.20	62	0.41	41.79	62	0.25	27.93
HNRPC_HUMAN	Heterogeneous nuclear ribonucleoproteins C1/C2	cellular	33.6	4.8	217	8	21.2	20	1.48	30.81	20	0.42	34.26	20	0.28	42.84
HMGB1_HUMAN	High mobility group protein B1 (High mobility group protein B1)	cellular	24.9	5.5	416	10	45.6	20	1.58	27.26	20	0.40	41.05	20	0.28	34.15
HINT1_HUMAN	Histidine triad nucleotide-binding protein 1	cellular	13.8	6.5	315	3	46.0	7	1.21	24.17	7	0.44	31.47	7	0.25	44.08
HINT2_HUMAN	Histidine triad nucleotide-binding protein 2	cellular	17.2	9.8	104	2	19.6	3	1.50	15.46	3	0.66	28.42	3	0.47	14.09
HRG_HUMAN	Histidine-rich glycoprotein precursor	blood	59.5	7.2	555	9	23.6	13	1.07	22.74	13	0.64	19.44	13	0.96	26.84
SAP18_HUMAN	Histone deacetylase complex subunit SAP18	cellular	17.5	9.9	74	3	24.2	2	1.47	45.49	2	1.04	15.08	2	0.51	42.00
H10_HUMAN	Histone H1.0 (Histone H1(0))	cellular	20.9	11.3	216	9	31.4	9	2.57	39.02	9	0.66	53.18	9	0.38	41.66
H12_HUMAN	Histone H1.2 (Histone H1d) - Homo sapiens	cellular	21.4	11.4	582	11	47.4	30	1.81	21.33	30	0.39	25.66	30	0.36	33.51
H13_HUMAN	Histone H1.3 (Histone H1c) - Homo sapiens	cellular	22.3	11.5	489	6	31.7	23	1.78	21.29	23	0.40	27.01	23	0.35	27.11
H14_HUMAN	Histone H1.4 (Histone H1b) - Homo sapiens	cellular	21.9	11.5	619	11	46.6	31	2.02	31.49	31	0.41	26.89	31	0.36	29.31
H15_HUMAN	Histone H1.5 (Histone H1a) - Homo sapiens	cellular	22.6	11.4	190	5	24.3	6	2.17	25.02	6	0.41	92.89	6	0.30	35.49
H1X_HUMAN	Histone H1x - Homo sapiens (Human)	cellular	22.5	11.2	52	1	9.9	3	1.45	5.56	3	0.58	11.28	3	0.59	36.48
H2A1H_HUMAN	Histone H2A type 1-H - Homo sapiens (Human)	cellular	13.9	11.3	430	6	46.9	31	2.13	27.90	31	0.45	33.01	30	0.29	25.99
H2A2B_HUMAN	Histone H2A type 2-B - Homo sapiens (Human)	cellular	14.0	11.3	305	5	40.8	25	2.09	27.38	25	0.42	21.81	25	0.27	36.61
H2AX_HUMAN	Histone H2A.x (H2a/x) - Homo sapiens (Human)	cellular	15.1	11.2	381	5	38.5	28	2.18	25.65	28	0.48	32.21	28	0.30	36.38
H2AV_HUMAN	Histone H2AV (H2A.F/Z) - Homo sapiens (Human)	cellular	13.5	11.0	295	4	31.2	25	1.93	38.86	25	0.53	26.79	25	0.36	35.23
H2B2F_HUMAN	Histone H2B type 2-F - Homo sapiens (Human)	cellular	13.9	10.8	487	9	63.5	48	2.27	44.81	48	0.55	24.96	47	0.33	34.95
H32_HUMAN	Histone H3.2 (H3/m) (H3/o) - Homo sapiens	cellular	15.4	11.9	230	7	28.7	42	2.31	26.95	42	0.47	23.12	41	0.57	34.40
H4_HUMAN	Histone H4 - Homo sapiens (Human)	cellular	11.4	11.8	746	9	59.2	70	2.15	41.53	70	0.44	24.87	70	0.41	42.16
I1C12_HUMAN	HLA class I histocompatibility antigen, Cw-12 alpha	cellular	40.9	5.9	151	3	10.7	4	1.23	18.71	4	0.72	28.29	4	0.25	25.47
2DRA_HUMAN	HLA class II histocompatibility antigen, DR alpha	cellular	28.6	4.8	65	2	9.4	2	1.66	32.50	2	0.72	1.26	2	0.35	13.87
F10A1_HUMAN	Hsc70-interacting protein (Hip) (Suppression of tumor progression)	cellular	41.3	5.0	88	3	9.5	3	1.31	12.64	3	0.82	38.86	3	0.32	36.72
IGHA1_HUMAN	Ig alpha-1 chain C region - Homo sapiens	blood	37.6	6.1	929	13	39.7	80	0.90	19.47	82	1.69	33.05	81	1.25	43.94
IGHA2_HUMAN	Ig alpha-2 chain C region - Homo sapiens	blood	36.5	5.7	844	11	35.6	74	0.88	18.44	76	1.65	42.42	75	1.11	49.79
IGHG1_HUMAN	Ig gamma-1 chain C region - Homo sapiens	blood	36.1	9.4	762	15	52.4	56	1.17	25.21	56	1.21	36.21	56	0.78	29.53
IGHG2_HUMAN	Ig gamma-2 chain C region - Homo sapiens	blood	35.9	8.8	697	9	40.2	36	1.03	11.49	36	1.72	37.18	36	1.16	30.85
IGHG3_HUMAN	Ig gamma-3 chain C region (Heavy chain disease)	blood	32.3	9.1	504	11	37.2	34	1.16	18.61	34	1.21	39.42	34	0.88	28.69
IGHG4_HUMAN	Ig gamma-4 chain C region - Homo sapiens	blood	35.9	7.8	449	7	30.6	19	1.00	7.76	19	1.40	26.36	19	0.91	39.04
HV305_HUMAN	Ig heavy chain V-III region BRO - Homo sapiens	blood	13.2	7.0	180	2	29.2	8	0.97	6.35	8	1.05	30.66	8	0.83	23.12
HV315_HUMAN	Ig heavy chain V-III region WAS - Homo sapiens	blood	13.1	9.2	85	1	41.0	4	1.15	17.13	4	1.17	10.21	4	0.70	35.06
KAC_HUMAN	Ig kappa chain C region - Homo sapiens	blood	11.6	5.5	725	7	86.8	46	0.99	25.76	46	0.94	26.47	46	1.07	21.56
KV101_HUMAN	Ig kappa chain V-I region AG - Homo sapiens	blood	12.0	5.6	344	2	31.5	9	1.15	16.12	9	0.73	23.82	9	0.69	25.84
KV121_HUMAN	Ig kappa chain V-I region Ni - Homo sapiens	blood	12.2	5.1	170	1	30.4	4	1.21	19.62	4	0.69	7.41	4	0.73	22.81
KV118_HUMAN	Ig kappa chain V-I region WEA - Homo sapiens	blood	11.8	9.6	316	3	36.1	7	1.15	22.57	7	0.81	27.88	7	0.68	23.86
KV203_HUMAN	Ig kappa chain V-II region MIL - Homo sapiens	blood	12.0	0.0	128	2	21.4	12	1.16	15.06	12	0.82	21.21	12	0.82	36.23
KV204_HUMAN	Ig kappa chain V-II region TEW - Homo sapiens	blood	12.3	5.6	271	2	38.9	14	1.11	21.28	14	0.84	20.87	14	0.80	35.64
KV302_HUMAN	Ig kappa chain V-III region SIE - Homo sapiens	blood	11.8	9.5	277	3	45.9	22	1.12	23.13	22	0.75	28.76	22	0.70	27.95
KV309_HUMAN	Ig kappa chain V-III region VG precursor	blood	12.6	4.7	64	1	26.1	10	1.10	17.55	10	0.69	36.86	10	0.54	36.05
KV310_HUMAN	Ig kappa chain V-III region VH precursor	blood	12.7	5.6	82	2	23.3	6	1.40	10.23	6	0.96	12.98	6	0.55	29.27
KV402_HUMAN	Ig kappa chain V-IV region Len - Homo sapiens	blood	12.6	9.0	304	3	36.8	8	1.05	28.17	8	0.83	23.76	8	0.57	39.93
LAC_HUMAN	Ig lambda chain C regions - Homo sapiens	blood	11.2	7.6	497	6	75.2	50	0.98	16.19	50	0.72	23.57	50	1.07	25.37
LV102_HUMAN	Ig lambda chain V-I region HA - Homo sapiens	blood	11.9	9.7	117	3	27.7	4	0.91	14.48	4	0.83	33.60	4	0.89	6.40
MUC_HUMAN	Ig mu chain C region - Homo sapiens	blood	49.5	6.4	328	8	22.9	14	1.12	25.64	14	1.33	41.92	14	1.09	45.26
IGJ_HUMAN	Immunoglobulin J chain - Homo sapiens	blood	15.6	4.4	464	6	48.2	29	1.05	15.66	29	0.76	17.82	29	1.16	43.67
IPYR_HUMAN	Inorganic pyrophosphatase (EC 3.6.1.1) (Pyrophosphatase)	cellular	32.6	5.5	181	5	15.9	12	1.32	16.37	12	0.59	14.06	12	0.24	26.08
ILF3_HUMAN	Interleukin enhancer-binding factor 3 (Nuclear factor kappa B)	cellular	95.3	9.3	84	2	3.4	2	2.14	34.99	2	0.46	10.83	2	0.29	4.46

IDHP_HUMAN	Isocitrate dehydrogenase [NADP], mitochondrial	cellular	50.9	9.6	158	6	12.8	8	1.46	19.42	8	0.61	25.52	8	0.26	51.03
K1C15_HUMAN	Keratin, type I cytoskeletal 15 (Cytokeratin-15)	cellular	49.1	4.6	424	6	16.0	27	1.51	19.12	26	0.52	36.09	27	0.38	40.07
K1C18_HUMAN	Keratin, type I cytoskeletal 18 (Cytokeratin-18)	cellular	48.0	5.2	408	11	23.5	29	1.58	28.51	29	0.56	23.06	29	0.49	25.64
K1C19_HUMAN	Keratin, type I cytoskeletal 19 (Cytokeratin-19)	cellular	44.1	4.9	1419	27	59.2	89	1.64	30.65	88	0.38	45.60	89	0.28	39.16
K1C9_HUMAN	Keratin, type I cytoskeletal 9 (Cytokeratin-9)	cellular	62.1	5.1	169	3	10.9	6	1.48	14.49	6	0.59	71.36	6	0.41	39.41
K2C1_HUMAN	Keratin, type II cytoskeletal 1 (Cytokeratin-1)	cellular	66.0	8.8	238	4	15.1	11	2.42	48.25	11	0.39	73.37	11	0.41	86.34
K22E_HUMAN	Keratin, type II cytoskeletal 2 epidermal (Cytokeratin-2)	cellular	65.8	8.9	306	2	10.9	10	2.84	32.36	10	0.22	24.97	10	0.22	36.84
K2C6E_HUMAN	Keratin, type II cytoskeletal 6E (Cytokeratin-6E)	cellular	60.0	8.9	366	6	16.7	31	2.22	40.26	31	0.44	59.49	31	0.35	45.91
K2C7_HUMAN	Keratin, type II cytoskeletal 7 (Cytokeratin-7)	cellular	51.4	5.4	1304	26	45.6	99	1.81	40.06	99	0.53	39.29	98	0.27	40.59
K2C8_HUMAN	Keratin, type II cytoskeletal 8 (Cytokeratin-8)	cellular	53.7	5.4	1238	25	49.3	87	2.19	32.16	87	0.36	46.17	87	0.36	36.08
KTN1_HUMAN	Kinectin (Kinesin receptor) (CG-1 antigen) - Homo sapiens	cellular	156.2	5.4	144	5	3.8	7	1.65	27.96	7	0.79	33.68	7	0.55	41.07
TRFL_HUMAN	Lactotransferrin precursor (EC 3.4.21.-) (Lactoferrin)	extracellular	78.1	9.6	223	4	10.7	14	1.07	11.76	14	0.87	21.77	14	0.91	39.48
LGUL_HUMAN	Lactoylglycyl-L-histidine lyase (EC 4.4.1.5) (Methylglyoxal lyase)	cellular	20.8	5.0	140	3	21.7	8	2.02	25.75	8	0.56	20.11	8	0.45	33.41
LMNA_HUMAN	Lamin A/C (70 kDa lamin) (Renal carcinoma antigen)	cellular	74.1	6.6	2050	40	58.7	108	1.50	21.73	108	0.61	21.61	106	0.44	24.44
LAP2B_HUMAN	Lamina-associated polypeptide 2, isoforms beta	cellular	50.6	9.8	125	3	9.5	3	1.46	15.36	3	0.56	36.60	3	0.37	36.25
LMNB1_HUMAN	Lamin-B1 - Homo sapiens (Human)	cellular	66.4	5.0	187	5	13.7	8	1.69	9.40	8	0.54	11.40	8	0.36	26.06
LMNB2_HUMAN	Lamin-B2 - Homo sapiens (Human)	cellular	67.6	5.2	324	10	17.3	15	1.82	24.11	15	0.58	16.01	15	0.34	31.79
LAMB2_HUMAN	Laminin subunit beta-2 precursor (S-laminin) (Laminin)	extracellular	196.0	6.1	81	3	2.4	3	0.96	18.69	3	1.90	24.08	3	0.83	17.43
A2GL_HUMAN	Leucine-rich alpha-2-glycoprotein precursor (LRP)	blood	38.2	6.5	230	6	21.9	13	1.20	26.17	13	0.65	54.23	13	0.75	59.83
LRC59_HUMAN	Leucine-rich repeat-containing protein 59 - Homo sapiens	cellular	34.9	10.4	360	6	27.4	18	1.27	11.67	18	0.67	34.40	18	0.20	29.88
LASP1_HUMAN	LIM and SH3 domain protein 1 (LASP-1) (MLN 52)	cellular	29.7	6.7	273	8	31.4	11	1.49	20.81	11	0.49	26.83	11	0.32	19.64
LPP_HUMAN	Lipoma-preferred partner (LIM domain-containing protein)	cellular	65.7	7.8	177	5	7.8	7	1.83	23.24	7	0.61	31.16	7	0.46	26.82
EST1_HUMAN	Liver carboxylesterase 1 precursor (EC 3.1.1.1)	cellular	62.5	6.2	108	3	5.6	3	1.23	43.36	3	1.41	13.84	3	0.72	45.37
LDHA_HUMAN	L-lactate dehydrogenase A chain (EC 1.1.1.27)	cellular	36.7	9.3	48	2	5.7	2	1.08	9.55	2	0.72	4.15	2	0.43	19.26
ACSL1_HUMAN	Long-chain-fatty-acid--CoA ligase 1 (EC 6.2.1.3)	cellular	77.9	7.0	87	2	5.9	2	0.99	8.51	2	1.17	14.55	2	0.99	19.12
LUM_HUMAN	Lumican precursor (Keratan sulfate proteoglycan core protein)	extracellular	38.4	6.2	1216	17	47.3	112	1.04	13.60	111	0.63	36.44	111	0.87	29.20
LA_HUMAN	Lupus La protein (Sjogren syndrome type B antigen)	cellular	46.8	6.8	48	2	4.7	7	1.91	19.88	7	0.32	28.84	6	0.35	25.79
LAMP2_HUMAN	Lysosome-associated membrane glycoprotein 2	cellular	44.9	5.3	34	1	2.0	6	1.52	17.22	6	0.59	19.02	6	0.35	31.93
MIF_HUMAN	Macrophage migration inhibitory factor (MIF)	extracellular	12.5	9.1	157	2	17.4	13	1.92	40.58	13	0.61	27.52	13	0.19	49.34
CAPG_HUMAN	Macrophage-capping protein (Actin-regulatory protein)	cellular	38.5	5.9	245	5	21.0	15	1.30	30.44	15	0.51	31.31	15	0.36	39.61
MDHC_HUMAN	Malate dehydrogenase, cytoplasmic (EC 1.1.1.3)	cellular	36.4	7.7	131	3	8.7	9	1.17	15.41	9	0.93	27.06	9	0.58	14.39
MDHM_HUMAN	Malate dehydrogenase, mitochondrial precursor	cellular	35.5	9.8	260	5	19.5	24	1.13	30.63	24	0.52	38.65	24	0.47	46.93
CBPA3_HUMAN	Mast cell carboxypeptidase A precursor (EC 3.4.11.1)	cellular	48.7	9.6	125	4	9.6	11	1.53	19.72	11	0.83	18.65	11	0.70	27.41
AOC3_HUMAN	Membrane copper amine oxidase (EC 1.4.3.6)	cellular	84.6	6.1	249	6	11.3	10	1.19	23.15	10	1.05	90.90	10	0.44	78.49
PGRC1_HUMAN	Membrane-associated progesterone receptor component 1	cellular	21.7	4.4	104	4	19.5	12	1.42	21.40	12	0.70	13.62	12	0.34	24.17
PGRC2_HUMAN	Membrane-associated progesterone receptor component 2	cellular	23.8	4.6	129	3	21.1	7	1.31	12.27	8	0.75	19.00	8	0.33	17.71
MFAP2_HUMAN	Microfibrillar-associated protein 2 precursor (Mfap2)	extracellular	20.8	4.7	54	2	10.4	3	1.36	8.47	3	0.63	6.60	3	0.26	42.12
MFAP5_HUMAN	Microfibrillar-associated protein 5 precursor (Mfap5)	extracellular	19.6	5.6	89	3	20.8	3	1.52	27.73	3	1.27	13.53	3	0.34	28.67
MARE1_HUMAN	Microtubule-associated protein RP/EB family member 1	cellular	30.0	4.9	114	2	10.8	3	1.64	16.67	3	0.62	27.42	3	0.30	11.05
MIME_HUMAN	Mimecan precursor (Osteoglycin) (Osteonidogen)	extracellular	33.9	5.3	1050	17	47.0	66	1.09	14.96	66	1.06	26.34	65	0.88	35.13
MOES_HUMAN	Moesin (Membrane-organizing extension spike protein)	cellular	67.8	6.0	288	6	12.8	18	1.47	26.63	18	0.78	17.87	18	0.38	21.46
MUC5B_HUMAN	Mucin-5B precursor (Mucin-5 subtype B, tracheobronchial)	extracellular	590.1	6.2	295	4	1.2	12	1.24	38.47	12	0.18	66.99	12	0.65	36.10
MYL6_HUMAN	Myosin light polypeptide 6 (Smooth muscle and nonmuscle)	cellular	16.9	4.4	532	10	56.3	40	1.39	28.91	40	0.44	44.66	40	0.24	44.89
MLRM_HUMAN	Myosin regulatory light chain 2, nonsarcomeric	cellular	19.8	4.5	392	6	39.2	27	1.59	19.35	27	0.48	21.71	27	0.20	33.51
MLRN_HUMAN	Myosin regulatory light chain 2, smooth muscle	cellular	19.8	4.6	386	5	39.0	20	1.65	20.60	20	0.50	26.31	20	0.28	38.11
MYH9_HUMAN	Myosin-9 (Myosin heavy chain 9) (Myosin heavy chain)	cellular	226.4	5.4	2048	40	25.9	60	1.43	19.13	60	0.66	27.85	60	0.29	39.05
NEDD8_HUMAN	NEDD8 precursor (Ubiquitin-like protein Nedd8)	cellular	9.1	9.1	48	1	25.9	3	1.26	28.53	3	0.54	27.43	3	0.25	29.40
AHNK_HUMAN	Neuroblast differentiation-associated protein A	cellular	312.3	6.3	2052	38	15.0	77	1.54	27.73	77	0.63	24.26	77	0.44	27.76
DEF1_HUMAN	Neutrophil defensin 1 precursor (HNP-1) (HP-1)	extracellular	10.2	7.6	84	3	25.5	6	1.48	18.28	6	0.95	13.32	6	0.52	14.65
DDAH2_HUMAN	NG,NG-dimethylarginine dimethylaminohydrolase	cellular	29.6	5.6	115	4	14.0	5	2.16	35.80	5	0.72	31.84	5	0.51	39.43
NID2_HUMAN	Nidogen-2 precursor (NID-2) (Osteonidogen) - Homo sapiens	extracellular	151.3	5.0	199	5	4.7	7	1.45	13.23	7	1.35	26.62	7	0.70	30.72
NGBR_HUMAN	Nogo-B receptor precursor (NgBR) (Nuclear undifferentiated protein)	cellular	33.2	9.9	35	1	2.4	4	1.66	6.25	4	0.46	26.28	4	0.23	2.11
HMGN2_HUMAN	Nonhistone chromosomal protein HMG-17 (High mobility group protein N2)	cellular	9.4	10.5	154	3	47.8	14	1.47	27.06	14	0.33	24.71	14	0.21	26.88
NLTP_HUMAN	Nonspecific lipid-transfer protein (EC 2.3.1.176)	cellular	59.0	6.5	99	3	2.6	4	1.91	37.13	4	0.34	63.09	4	0.22	10.86
NUCB1_HUMAN	Nucleobindin-1 precursor (CALNUC) - Homo sapiens	cellular	53.8	5.0	120	4	12.8	4	1.22	7.92	4	0.51	15.07	4	0.35	30.64
NUCL_HUMAN	Nucleolin (Protein C23) - Homo sapiens (Human)	cellular	76.6	4.4	638	18	25.9	31	1.30	25.96	31	0.44	30.51	31	0.25	33.92
NPM_HUMAN	Nucleophosmin (NPM) (Nucleolar phosphoprotein)	cellular	32.6	4.5	345	10	27.2	15	1.55	22.77	15	0.45	42.58	15	0.27	32.00

NDBK_HUMAN	Nucleoside diphosphate kinase B (EC 2.7.4.6) (Nucleoside diphosphate kinase B)	cellular	17.3	9.4	365	8	46.1	36	1.54	20.86	36	0.52	30.05	36	0.13	38.23
NP1L1_HUMAN	Nucleosome assembly protein 1-like 1 (NAP-1-like 1)	cellular	45.3	4.2	63	3	10.2	3	1.70	65.88	3	0.58	24.82	3	0.37	57.89
PTMS_HUMAN	Parathymosin - Homo sapiens (Human)	cellular	11.5	4.0	114	3	23.5	9	1.78	30.53	9	0.51	28.17	9	0.35	38.49
PDL1_HUMAN	PDZ and LIM domain protein 1 (Elfin) (LIM domain protein 1)	cellular	36.0	6.6	44	2	8.8	7	1.26	30.44	7	0.75	21.65	7	0.42	20.46
PDLI7_HUMAN	PDZ and LIM domain protein 7 (LIM domain protein 7)	cellular	49.8	9.9	73	2	5.9	7	1.58	15.59	7	0.63	32.27	6	0.35	24.20
PPIA_HUMAN	Peptidyl-prolyl cis-trans isomerase A (EC 5.2.1.8)	cellular	18.0	9.0	934	16	69.1	86	1.61	28.15	86	0.40	27.46	86	0.24	37.45
PPIB_HUMAN	Peptidyl-prolyl cis-trans isomerase B precursor	cellular	22.7	9.9	439	12	49.0	25	1.32	26.52	25	0.47	37.59	25	0.24	37.67
PLIN_HUMAN	Perilipin (PERI) (Lipid droplet-associated protein)	cellular	55.9	6.0	633	11	29.7	31	1.06	30.05	31	1.20	33.65	31	0.75	30.29
POSTN_HUMAN	Periostin precursor (PN) (Osteoblast-specific factor)	extracellular	93.3	7.8	920	11	24.9	36	1.36	11.13	36	0.47	16.46	36	0.28	32.63
PRDX1_HUMAN	Peroxiredoxin-1 (EC 1.11.1.15) (Thioredoxin peroxidase)	cellular	22.1	9.2	881	14	60.8	67	1.54	29.57	67	0.53	24.80	65	0.19	37.51
PRDX2_HUMAN	Peroxiredoxin-2 (EC 1.11.1.15) (Thioredoxin peroxidase)	cellular	21.9	5.6	463	10	54.5	49	1.59	27.69	49	0.68	22.19	49	0.35	36.53
PRDX5_HUMAN	Peroxiredoxin-5, mitochondrial precursor (EC 1.11.1.15)	cellular	22.0	9.9	358	6	36.0	10	1.45	15.04	10	0.47	28.20	10	0.24	32.74
PRDX6_HUMAN	Peroxiredoxin-6 (EC 1.11.1.15) (Antioxidant peroxidase)	cellular	25.0	6.0	663	11	58.5	38	1.43	34.43	38	0.80	23.74	37	0.52	28.97
PEBP1_HUMAN	Phosphatidylethanolamine-binding protein 1 (PEBP1)	cellular	21.0	7.8	546	12	60.4	35	1.74	38.19	35	0.49	26.76	35	0.38	35.92
PGK1_HUMAN	Phosphoglycerate kinase 1 (EC 2.7.2.3) (Primer kinase)	cellular	44.6	9.2	757	18	43.2	47	1.55	28.69	47	0.58	27.03	47	0.26	41.54
PGAM1_HUMAN	Phosphoglycerate mutase 1 (EC 5.4.2.1) (EC 5.4.2.1)	cellular	28.8	6.8	109	3	19.7	3	1.83	13.09	3	1.55	21.20	3	0.52	16.84
PEDF_HUMAN	Pigment epithelium-derived factor precursor (PEDF)	cellular	46.3	6.0	75	3	6.2	3	1.15	4.44	3	0.76	10.31	3	0.66	12.73
IC1_HUMAN	Plasma protease C1 inhibitor precursor (C1 inh)	blood	55.1	6.1	179	6	14.6	7	1.28	13.89	7	0.82	16.74	7	0.92	33.80
PLMN_HUMAN	Plasminogen precursor (EC 3.4.21.7) [Contains: plasminogen]	blood	90.5	7.3	85	3	4.7	5	1.30	13.11	5	0.85	8.42	5	0.67	22.58
PLEC1_HUMAN	Plectin-1 (PLTN) (PCN) (Hemidesmosomal protein)	cellular	531.4	5.6	804	20	7.4	21	1.21	39.71	22	0.63	28.35	21	0.36	31.37
PCBP1_HUMAN	Poly(rC)-binding protein 1 (Alpha-CP1) (hnRNP-C1)	cellular	37.5	6.8	123	4	19.4	4	1.14	36.74	4	0.85	23.33	4	0.40	27.05
PABP1_HUMAN	Polyadenylate-binding protein 1 (Poly(A)-binding protein 1)	cellular	70.6	10.0	102	4	8.3	5	1.47	43.94	5	0.69	35.65	5	0.33	10.97
PTRF_HUMAN	Polymerase I and transcript release factor - Homo sapiens (Human)	cellular	43.4	5.4	667	12	36.7	18	1.37	24.16	18	1.07	28.02	18	1.09	34.17
PIGR_HUMAN	Polymeric-immunoglobulin receptor precursor	cellular	83.3	5.5	73	1	3.7	2	1.24	14.85	2	3.44	3.76	2	1.11	11.53
PFD2_HUMAN	Prefoldin subunit 2 - Homo sapiens (Human)	cellular	16.6	6.2	81	1	23.4	3	1.57	12.73	3	0.61	26.85	3	0.32	10.54
SAP_HUMAN	Proactivator polypeptide precursor [Contains: proactivator]	cellular	58.1	4.9	302	6	11.8	29	1.42	27.13	28	0.34	26.21	28	0.20	32.29
DDX17_HUMAN	Probable ATP-dependent RNA helicase DDX17 (DDX17)	cellular	72.3	9.5	222	7	16.2	13	1.45	20.99	13	0.60	17.34	13	0.41	29.20
PCOC1_HUMAN	Procollagen C-endopeptidase enhancer 1 precursor	cellular	47.9	8.7	87	3	15.4	5	1.47	30.33	4	0.71	17.50	4	0.42	30.45
PROF1_HUMAN	Profilin-1 (Profilin I) - Homo sapiens (Human)	cellular	15.0	9.4	727	10	61.4	29	1.79	33.15	29	0.53	28.02	29	0.28	30.96
PDCD6_HUMAN	Programmed cell death protein 6 (Probable caspase-3 activator)	cellular	21.9	5.0	77	2	8.9	2	0.99	24.24	2	0.42	6.48	2	0.34	19.46
PIP_HUMAN	Prolactin-inducible protein precursor (Prolactin-inducible protein)	extracellular	16.6	9.3	310	5	65.1	19	1.17	24.98	19	1.11	48.48	19	0.56	38.30
PRELP_HUMAN	Prolargin precursor (Proline-arginine-rich end leucine-rich repeat-containing protein)	extracellular	43.8	10.2	821	14	42.4	50	1.20	19.99	50	0.93	31.12	50	0.58	24.70
PSME1_HUMAN	Proteasome activator complex subunit 1 (Proteasome activator complex subunit 1)	cellular	28.7	5.7	364	7	36.9	11	1.49	19.69	12	0.46	35.62	11	0.25	27.09
CN166_HUMAN	Protein C14orf166 - Homo sapiens (Human)	cellular	28.1	6.2	40	1	3.3	3	1.33	11.31	3	0.88	16.52	3	0.41	6.73
CUTA_HUMAN	Protein CutA precursor (Brain acetylcholinesterase)	cellular	19.1	5.3	66	1	7.8	5	1.48	13.15	5	0.81	16.43	5	0.63	21.73
PDIA3_HUMAN	Protein disulfide-isomerase A3 precursor (EC 5.0.2.1)	cellular	56.7	5.9	1143	25	43.2	40	1.48	24.54	40	0.43	32.56	40	0.27	41.04
PDIA4_HUMAN	Protein disulfide-isomerase A4 precursor (EC 5.0.2.1)	cellular	72.9	4.8	117	5	8.7	6	1.36	31.56	6	0.46	16.66	6	0.36	38.62
PDIA6_HUMAN	Protein disulfide-isomerase A6 precursor (EC 5.0.2.1)	cellular	48.1	4.8	176	3	12.3	6	1.31	26.78	6	0.54	17.65	6	0.26	57.56
PDIA1_HUMAN	Protein disulfide-isomerase precursor (EC 5.0.2.1)	cellular	57.1	4.6	1014	18	41.1	47	1.44	24.10	47	0.36	18.49	47	0.18	28.51
PARK7_HUMAN	Protein DJ-1 (Oncogene DJ1) (Parkinson disease protein)	cellular	19.9	6.4	361	7	50.8	12	1.35	21.03	12	0.58	23.38	12	0.33	22.99
S10AA_HUMAN	Protein S100-A10 (S100 calcium-binding protein S100-A10)	cellular	11.2	7.7	146	4	37.1	7	1.29	19.89	7	0.40	26.47	7	0.38	21.37
S10AB_HUMAN	Protein S100-A11 (S100 calcium-binding protein S100-A11)	cellular	11.7	7.5	227	6	41.0	18	1.33	21.84	18	0.41	19.28	18	0.24	29.21
S10AD_HUMAN	Protein S100-A13 (S100 calcium-binding protein S100-A13)	cellular	11.5	5.9	122	2	23.5	6	1.67	10.95	6	0.41	43.53	6	0.41	14.40
S10A4_HUMAN	Protein S100-A4 (S100 calcium-binding protein S100-A4)	cellular	11.7	5.8	284	6	46.5	16	1.34	15.48	16	0.69	17.10	16	0.35	30.61
S10A6_HUMAN	Protein S100-A6 (S100 calcium-binding protein S100-A6)	cellular	10.2	5.2	247	5	41.1	32	1.40	15.36	32	0.50	27.32	32	0.19	25.88
S10A8_HUMAN	Protein S100-A8 (S100 calcium-binding protein S100-A8)	cellular	10.8	6.6	327	6	47.3	21	1.13	30.36	21	0.53	38.36	21	0.13	48.30
S10A9_HUMAN	Protein S100-A9 (S100 calcium-binding protein S100-A9)	cellular	13.2	5.7	407	7	63.2	13	1.40	22.71	13	0.53	38.01	13	0.11	32.51
SET_HUMAN	Protein SET (Phosphatase 2A inhibitor I2PP2A) (SET)	cellular	33.5	4.1	109	2	8.3	3	1.37	7.31	3	0.39	25.03	3	0.29	12.68
THRB_HUMAN	Prothrombin precursor (EC 3.4.21.5) (Coagulation factor II)	blood	70.0	5.6	553	12	30.1	19	1.02	28.12	19	0.80	37.69	19	0.84	25.50
PTMA_HUMAN	Prothymosin alpha [Contains: Thymosin alpha-1]	cellular	12.2	3.5	182	5	24.3	11	1.69	30.87	11	0.49	26.68	11	0.31	42.20
LC7L2_HUMAN	Putative RNA-binding protein Luc7-like 2 - Homo sapiens (Human)	cellular	46.5	10.8	118	3	11.0	4	1.82	24.66	4	0.77	30.34	4	0.32	15.07
PNPO_HUMAN	Pyridoxine-5'-phosphate oxidase (EC 1.4.3.5) (Pyridoxine-5'-phosphate oxidase)	cellular	30.0	6.7	78	2	11.9	2	1.84	24.21	2	0.30	17.59	2	0.21	62.24
KPYM_HUMAN	Pyruvate kinase isozymes M1/M2 (EC 2.7.1.40)	cellular	57.9	9.0	190	3	8.3	8	1.49	17.05	8	0.66	19.20	8	0.40	22.84
G3BP1_HUMAN	Ras GTPase-activating protein-binding protein 1 (G3BP1)	cellular	52.1	5.3	111	3	9.9	4	0.98	17.28	4	0.65	22.50	4	0.41	19.46
IQGA1_HUMAN	Ras GTPase-activating-like protein IQGAP1 (p190)	cellular	189.1	6.1	93	2	2.5	3	1.13	19.08	3	0.58	23.10	3	0.42	28.04
RAC2_HUMAN	Ras-related C3 botulinum toxin substrate 2 (Rac2)	cellular	21.4	8.9	56	2	10.4	3	1.46	26.48	3	0.60	27.00	3	0.30	31.24
RAB7A_HUMAN	Ras-related protein Rab-7a - Homo sapiens (Human)	cellular	23.5	7.2	107	3	19.3	5	1.27	19.30	5	0.84	15.27	5	0.46	12.22

RAP1B_HUMAN	Ras-related protein Rap-1b precursor (GTP-binding protein)	cellular	20.8	5.5	96	3	15.8	3	1.47	34.61	3	0.86	23.77	3	0.36	5.99
RCN1_HUMAN	Reticulocalbin-1 precursor - Homo sapiens (Human)	cellular	38.9	4.7	98	4	14.2	5	1.30	17.63	5	0.55	26.68	5	0.39	34.81
RCN3_HUMAN	Reticulocalbin-3 precursor (EF-hand calcium-binding protein)	cellular	37.5	4.6	112	5	26.5	4	0.93	25.97	5	0.33	43.29	4	0.33	22.92
RRBP1_HUMAN	Ribosome-binding protein 1 (Ribosome receptor)	cellular	152.4	9.3	455	13	12.9	15	1.09	24.05	15	0.49	31.30	15	0.26	34.34
SFRP1_HUMAN	Secreted frizzled-related protein 1 precursor (Secreted frizzled-related protein)	extracellular	35.4	10.1	42	1	8.3	6	1.76	26.17	6	1.28	43.06	6	0.94	61.31
SG1D2_HUMAN	Secretoglobin family 1D member 2 precursor (Secretoglobin)	extracellular	9.9	9.9	104	2	22.2	4	1.43	9.11	4	0.39	16.72	4	0.85	22.16
SBP1_HUMAN	Selenium-binding protein 1 - Homo sapiens (Human)	cellular	52.3	6.1	53	2	5.1	2	1.27	15.48	2	1.16	36.33	2	0.69	39.19
PAK2_HUMAN	Serine/threonine-protein kinase PAK 2 (EC 2.7.11.1)	cellular	58.0	5.6	110	2	5.0	2	1.31	2.94	2	0.62	36.31	2	0.28	1.52
TRFE_HUMAN	Serotransferrin precursor (Transferrin) (Siderophilin)	blood	77.0	7.0	2372	42	59.7	163	0.94	18.58	165	1.05	28.35	162	0.89	29.44
SERPH_HUMAN	Serpin H1 precursor (Collagen-binding protein)	cellular	46.4	9.3	126	3	8.4	3	1.02	12.22	3	0.58	38.41	3	0.37	54.83
ALBU_HUMAN	Serum albumin precursor - Homo sapiens (Human)	plasma	69.3	5.9	5099	69	87.4	575	0.69	21.98	579	1.30	40.12	579	1.50	43.95
SH3L1_HUMAN	SH3 domain-binding glutamic acid-rich-like protein	cellular	12.8	5.1	325	6	56.1	10	1.56	24.80	10	0.34	30.10	10	0.27	32.45
SRP09_HUMAN	Signal recognition particle 9 kDa protein (SRP9)	cellular	10.1	9.1	72	2	25.6	2	1.33	14.86	2	0.48	22.51	2	0.32	11.52
SMAP_HUMAN	Small acidic protein - Homo sapiens (Human)	cellular	20.3	4.4	73	2	8.2	2	2.17	0.89	2	0.38	15.36	2	0.33	56.04
MUCL_HUMAN	Small breast epithelial mucin precursor (Protein)	extracellular	9.0	4.2	57	2	8.9	3	0.91	18.30	3	0.41	5.84	3	0.21	38.39
SUMO4_HUMAN	Small ubiquitin-related modifier 4 precursor (SUMO4)	cellular	10.7	7.5	69	1	12.6	4	1.49	35.32	4	0.45	37.44	4	0.30	29.31
SORCN_HUMAN	Sorcin (22 kDa protein) (CP-22) (V19) - Homo sapiens	cellular	21.7	5.2	113	2	15.2	2	1.94	0.51	2	0.60	72.77	2	0.36	18.33
SPRC_HUMAN	SPARC precursor (Secreted protein acidic and rich in cysteine)	extracellular	34.6	4.6	125	4	19.1	4	1.49	18.09	4	0.64	23.40	4	0.44	15.25
SPTA2_HUMAN	Spectrin alpha chain, brain (Spectrin, non-erythrocytic)	cellular	284.4	5.1	583	16	9.3	20	1.54	30.45	20	0.83	21.06	20	0.52	32.02
SPTB2_HUMAN	Spectrin beta chain, brain 1 (Spectrin, non-erythrocytic)	cellular	274.4	5.3	320	9	5.8	11	1.48	21.41	11	0.88	23.42	11	0.60	36.58
SF3A1_HUMAN	Splicing factor 3 subunit 1 (Spliceosome-associated)	cellular	88.8	5.0	55	2	2.5	2	1.65	11.43	2	0.36	68.66	2	0.27	21.61
SF3B2_HUMAN	Splicing factor 3B subunit 2 (Spliceosome-associated)	cellular	97.6	5.4	78	3	3.1	3	1.19	38.37	3	0.62	21.60	3	0.30	27.01
SFRS1_HUMAN	Splicing factor, arginine/serine-rich 1 (pre-mRNA splicing)	cellular	27.7	10.8	283	9	44.4	14	1.19	26.06	14	0.60	37.67	14	0.24	23.92
SFRS2_HUMAN	Splicing factor, arginine/serine-rich 2 (Splicing factor)	cellular	25.5	12.4	80	2	9.5	9	1.74	26.02	9	0.33	24.29	9	0.20	50.49
SFRS3_HUMAN	Splicing factor, arginine/serine-rich 3 (Pre-mRNA splicing)	cellular	19.3	12.3	121	3	18.9	7	1.66	25.55	7	0.44	50.01	7	0.19	36.68
SFRS7_HUMAN	Splicing factor, arginine/serine-rich 7 (Splicing factor)	cellular	27.4	12.4	141	2	21.0	7	1.66	19.15	7	0.46	43.53	7	0.20	31.56
SFPO_HUMAN	Splicing factor, proline- and glutamine-rich (Polypyridine-binding)	cellular	76.1	9.9	119	4	10.6	4	1.59	10.52	4	0.41	26.34	4	0.27	42.45
SLIRP_HUMAN	SRA stem-loop-interacting RNA-binding protein	cellular	12.3	11.0	80	2	21.1	2	1.52	6.68	2	0.58	19.66	2	0.25	15.81
STMN1_HUMAN	Stathmin (Phosphoprotein p19) (pp19) (Oncoprotein)	cellular	17.3	5.7	181	5	26.8	12	1.07	33.76	12	0.79	39.47	12	0.20	30.12
GRP75_HUMAN	Stress-70 protein, mitochondrial precursor (75 kDa)	cellular	73.6	5.8	194	4	8.7	10	1.33	18.38	10	0.49	17.67	10	0.35	41.84
SUCB1_HUMAN	Succinyl-CoA ligase [ADP-forming] beta-chain, mitochondrial	cellular	50.3	7.7	69	1	6.5	4	0.54	9.63	4	1.03	20.32	4	1.18	8.53
SODC_HUMAN	Superoxide dismutase [Cu-Zn] (EC 1.15.1.1) - Homo sapiens	cellular	15.9	5.7	425	8	64.9	26	1.59	22.07	26	0.55	27.20	26	0.31	28.37
SODM_HUMAN	Superoxide dismutase [Mn], mitochondrial precursor	cellular	24.7	9.1	163	3	16.2	3	1.23	18.81	3	0.89	28.80	3	0.35	36.20
SDC1_HUMAN	Syndecan-1 precursor (SYND1) (CD138 antigen)	cellular	32.5	4.4	37	1	5.5	3	1.06	22.77	3	0.50	21.41	3	0.37	16.38
STX7_HUMAN	Syntaxin-7 - Homo sapiens (Human)	cellular	29.8	5.3	51	2	15.7	2	1.11	19.30	2	0.34	61.03	1	0.35	
TLN1_HUMAN	Talin-1 - Homo sapiens (Human)	cellular	269.6	5.7	441	9	8.1	18	1.20	25.42	18	0.65	28.83	18	0.43	19.51
TENX_HUMAN	Tenascin-X precursor (TN-X) (Hexabrachion-like protein)	extracellular	464.2	5.1	146	4	2.1	4	1.27	8.70	4	1.04	24.25	4	0.44	18.11
THIO_HUMAN	Thioredoxin (Trx) (ATL-derived factor) (ADF) (S)	cellular	11.7	4.7	231	6	59.0	10	1.31	29.35	10	0.49	40.81	10	0.29	41.76
TXD12_HUMAN	Thioredoxin domain-containing protein 12 precursor	cellular	19.2	5.1	61	2	14.0	3	1.84	25.71	3	0.89	10.30	3	0.52	15.13
TXND5_HUMAN	Thioredoxin domain-containing protein 5 precursor	cellular	47.6	5.6	201	7	13.4	10	1.40	21.60	10	0.45	27.29	10	0.26	47.15
PRDX3_HUMAN	Thioredoxin-dependent peroxide reductase, mitochondrial	cellular	27.7	8.9	124	3	10.5	5	1.35	25.03	5	0.53	15.86	5	0.35	38.55
THY1_HUMAN	Thy-1 membrane glycoprotein precursor (Thy-1)	cellular	17.9	9.8	78	2	18.0	2	1.48	18.42	2	0.59	13.68	2	0.40	52.91
TYB10_HUMAN	Thymosin beta-10 - Homo sapiens (Human)	cellular	5.0	5.2	178	6	47.7	20	1.51	29.95	20	0.28	48.82	20	0.13	36.53
TYB4_HUMAN	Thymosin beta-4 (T beta 4) (Fx) [Contains: Hemophilin]	cellular	5.0	4.9	152	4	45.5	21	1.70	27.73	21	0.22	33.12	21	0.15	40.38
TOIP1_HUMAN	Torsin-1A-interacting protein 1 - Homo sapiens	cellular	66.2	8.9	55	2	4.5	2	1.19	23.54	2	0.59	6.45	2	0.43	5.19
BGH3_HUMAN	Transforming growth factor-beta-induced protein	extracellular	74.6	8.6	313	4	11.4	8	0.93	17.63	8	1.03	18.67	8	0.69	37.42
RHOA_HUMAN	Transforming protein RhoA precursor (H12) - Homo sapiens	cellular	21.8	5.8	355	7	32.1	9	1.42	19.46	9	0.41	25.18	9	0.32	34.67
TAGL_HUMAN	Transgelin (Smooth muscle protein 22-alpha) (S)	cellular	22.6	9.4	655	15	60.2	43	1.60	25.49	43	0.65	19.60	43	0.36	21.10
TAGL2_HUMAN	Transgelin-2 (SM22-alpha homolog) - Homo sapiens	cellular	22.4	9.3	505	9	42.7	22	1.14	36.44	22	0.52	36.08	22	0.21	20.21
TERA_HUMAN	Transitional endoplasmic reticulum ATPase (TER)	cellular	89.3	5.0	225	6	11.2	8	1.40	25.10	8	0.66	19.23	8	0.44	8.77
TKT_HUMAN	Transketolase (EC 2.2.1.1) (TK) - Homo sapiens	cellular	67.8	8.5	128	3	6.4	5	1.31	22.77	5	1.38	17.39	5	0.30	29.38
TCTP_HUMAN	Translationally-controlled tumor protein (TCTP)	cellular	19.6	4.7	121	3	17.4	3	1.13	27.62	3	0.41	56.21	3	0.18	64.41
SSRA_HUMAN	Translocon-associated protein subunit alpha precursor	cellular	32.2	4.2	83	2	8.0	4	1.22	27.54	4	0.47	23.02	4	0.28	14.12
TTHY_HUMAN	Transthyretin precursor (Prealbumin) (TBPA) (T)	blood	15.9	5.4	672	6	62.6	30	1.01	14.08	30	0.66	25.37	30	1.42	28.49
TFF1_HUMAN	Trefoil factor 1 precursor (pS2 protein) (HP1.A)	extracellular	9.1	4.1	43	1	10.7	5	1.20	11.60	5	0.14	98.16	5	0.39	23.26
ECHA_HUMAN	Trifunctional enzyme subunit alpha, mitochondrial	cellular	82.9	9.8	101	3	8.0	3	1.66	15.91	4	0.24	277.69	3	0.35	19.06
TPIS_HUMAN	Triosephosphate isomerase (EC 5.3.1.1) (TIM) (T)	cellular	26.7	6.5	892	15	62.2	44	1.51	22.93	44	0.46	30.06	44	0.24	31.95

TPM3_HUMAN	Tropomyosin alpha-3 chain (Tropomyosin-3) (Tropomyosin-3)	cellular	32.8	4.5	648	17	39.1	37	1.55	25.24	37	0.42	37.50	37	0.24	38.23
TPM4_HUMAN	Tropomyosin alpha-4 chain (Tropomyosin-4) (Tropomyosin-4)	cellular	28.5	4.5	1058	25	66.9	59	1.53	24.96	59	0.41	35.04	59	0.24	36.52
TPM2_HUMAN	Tropomyosin beta chain (Tropomyosin 2) (Beta-tropomyosin)	cellular	32.8	4.5	937	20	49.6	45	1.61	23.55	45	0.40	33.45	45	0.24	34.94
TPM1_HUMAN	Tropomyosin-1 alpha chain (Alpha-tropomyosin)	cellular	32.7	4.5	887	22	51.1	38	1.46	28.83	38	0.48	34.56	38	0.30	37.01
TRY1_HUMAN	Trypsin-1 precursor (EC 3.4.21.4) (Trypsin I) (Cathepsin B)	extracellular	26.5	6.1	85	2	7.3	4	1.21	13.82	4	0.70	16.33	4	0.73	14.66
TRYA1_HUMAN	Trypsin alpha-1 precursor (EC 3.4.21.59) (Trypsin)	extracellular	30.8	6.6	199	3	22.2	5	1.25	14.43	5	1.02	22.48	5	0.97	25.33
TBA6_HUMAN	Tubulin alpha-6 chain (Alpha-tubulin 6) - Homo sapiens (Human)	cellular	49.9	4.8	639	12	36.7	20	1.38	24.99	20	0.91	32.19	20	0.37	43.19
TBAK_HUMAN	Tubulin alpha-ubiquitous chain (Alpha-tubulin)	cellular	50.1	4.8	653	12	36.6	22	1.40	24.46	22	0.94	30.41	22	0.35	42.83
TBB5_HUMAN	Tubulin beta chain (Tubulin beta-5 chain) - Homo sapiens (Human)	cellular	49.6	4.6	715	12	36.3	30	1.27	24.43	31	0.76	26.47	31	0.34	39.37
TBB2A_HUMAN	Tubulin beta-2A chain - Homo sapiens (Human)	cellular	49.9	4.6	578	9	30.8	28	1.27	24.64	28	0.75	22.13	28	0.33	37.67
TBB2C_HUMAN	Tubulin beta-2C chain (Tubulin beta-2 chain) - Homo sapiens (Human)	cellular	49.8	4.6	593	10	33.5	21	1.29	20.55	22	0.78	24.91	22	0.32	43.98
TPD54_HUMAN	Tumor protein D54 (hD54) (Tumor protein D52)	cellular	22.2	5.1	102	2	20.9	3	1.20	18.10	3	0.50	11.81	3	0.27	18.10
LSM3_HUMAN	U6 snRNA-associated Sm-like protein LSM3 - Homo sapiens (Human)	cellular	11.8	4.4	36	1	11.8	6	1.41	10.62	6	0.55	17.88	6	0.36	14.20
UCRH_HUMAN	Ubiquinol-cytochrome c reductase complex 11	cellular	10.7	4.2	78	3	35.2	3	1.06	14.63	3	0.72	14.24	3	0.40	41.76
UCR6_HUMAN	Ubiquinol-cytochrome c reductase complex 14	cellular	13.5	9.2	63	2	15.3	3	1.09	3.99	3	0.60	26.99	3	0.48	8.21
UBIQ_HUMAN	Ubiquitin - Homo sapiens (Human)	cellular	8.6	7.6	154	5	48.7	7	1.32	27.96	7	0.56	27.37	7	0.31	38.03
UBE1_HUMAN	Ubiquitin-activating enzyme E1 (A1S9 protein)	cellular	117.8	5.4	349	7	10.6	11	1.61	25.09	11	0.60	17.53	11	0.45	26.17
UB2L3_HUMAN	Ubiquitin-conjugating enzyme E2 L3 (EC 6.3.2.1)	cellular	17.9	9.5	179	5	45.5	7	1.19	14.31	7	0.56	27.56	7	0.27	13.08
UBE2N_HUMAN	Ubiquitin-conjugating enzyme E2 N (EC 6.3.2.19)	cellular	17.1	6.2	165	3	42.8	4	1.16	22.13	4	0.61	37.39	4	0.32	7.51
UFC1_HUMAN	Ufm1-conjugating enzyme 1 (Ubiquitin-fold modifier 1)	cellular	19.4	7.7	58	3	15.0	3	1.15	2.81	3	0.42	31.08	3	0.34	28.12
KCY_HUMAN	UMP-CMP kinase (EC 2.7.4.14) (Cytidylate kinase)	cellular	22.2	5.3	48	1	4.1	9	1.42	27.41	9	0.55	21.93	9	0.22	26.70
RD23B_HUMAN	UV excision repair protein RAD23 homolog B (hRAD23B)	cellular	43.1	4.6	110	2	4.6	2	1.71	40.71	2	0.58	10.82	2	0.33	10.08
CSPG2_HUMAN	Versican core protein precursor (Large fibroblast proteoglycan)	extracellular	372.6	4.3	351	7	3.4	21	1.17	24.85	21	0.36	36.18	21	0.54	27.33
VAPA_HUMAN	Vesicle-associated membrane protein-associated protein 1 (VAMP1)	cellular	27.9	9.6	74	2	8.0	3	1.55	7.16	3	0.64	33.79	3	0.40	28.60
VIME_HUMAN	Vimentin - Homo sapiens (Human)	cellular	53.6	4.9	3077	43	69.3	217	1.56	32.49	217	0.69	38.67	217	0.47	28.12
VINC_HUMAN	Vinculin (Metavinculin) - Homo sapiens (Human)	cellular	123.7	5.4	230	6	10.3	6	1.58	22.90	6	0.81	35.65	6	0.40	35.86
VTDB_HUMAN	Vitamin D-binding protein precursor (DBP) (GroEL)	extracellular	52.9	5.3	776	14	35.4	30	0.79	30.47	30	1.06	39.21	30	0.88	35.88
VTNC_HUMAN	Vitronectin precursor (Serum-spreading factor)	blood	54.3	5.5	321	7	16.1	14	1.30	21.08	14	0.75	17.25	14	0.68	18.82
ZA2G_HUMAN	Zinc-alpha-2-glycoprotein precursor (Zn-alpha-2-glycoprotein)	blood	33.9	5.5	902	14	53.6	51	1.38	27.01	51	0.67	32.27	51	0.96	41.55
ZYX_HUMAN	Zyxin (Zyxin-2) - Homo sapiens (Human)	cellular	61.2	6.2	80	2	5.1	4	1.19	25.72	4	0.59	16.29	4	0.34	39.20