Supplementary Table 1. Mass spectrometry-based quantation of UV-dependent HDAC8-Y100Bpa enrichment.

Accession	Description	Peptide count	Unique peptides	Ratio (+UV/-UV)	p value (+UV/-UV)	Pass Filters	UV 1 run 1	UV 1 run 2	UV 1 run 3	UV 2 run 1	UV 2 run 2	UV 2 run 3	UV 3 run 1	UV 3 run 2	UV 3 run 3	No UV 1 run 1	No UV 1 run 2	No UV 1 run 3	No UV 2 run 1	No UV 2 run 2	No UV 2 run 3	No UV 3 run 1	No UV 3 run 2	No UV 3 run 3
SYAC_HUMAN	Alanine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=AARS PE=1 SV=2	2	1	24.3	0.024	X	0.0	5.2	0.0	1769.6	3154.1	3808.8	168.8	2244.7	1568.4	0.0	0.0	0.0	251.5	64.5	198.5	8.6	0.0	0.0
PGAM4_HUMAN	Probable phosphoglycerate mutase 4 OS=Homo sapiens GN=PGAM4 PE=3 SV=1	5	1	7.3	0.027	X	192.7	44.9	51.4	17451.0	14046.1	10860.7	10419.7	1185.6	10632.5	321.5	166.1	5871.2	353.8	1568.5	321.9	70.3	138.6	117.6
SYEP_HUMAN	Bifunctional glutamate/proline--tRNA ligase OS=Homo sapiens GN=EPRS PE=1 SV=5	4	3	5.2	0.016	X	827.3	981.7	824.0	50925.0	56349.0	59492.3	28548.5	35192.6	40316.5	10921.6	14309.9	10766.5	4694.9	5011.7	5931.7	443.7	464.4	422.5
H14_HUMAN	Histone H1.4 OS=Homo sapiens GN=HIST1H1E PE=1 SV=2	5	1	5.0	0.017	X	922.7	136.8	830.3	7398.6	6133.8	12566.9	9450.9	8143.5	15530.8	2900.3	2254.5	2227.2	411.5	647.6	2499.4	533.4	328.1	500.7
K1C9_HUMAN	Keratin, type I cytoskeletal 9 OS=Homo sapiens GN=KRT9 PE=1 SV=3	30	24	4.8	0.015	X	31367.3	29764.8	30106.8	1730480.5	1562319.3	1395922.8	1893398.8	2379857.4	2099504.2	395402.3	477212.3	408308.6	309729.5	368570.2	333803.9	15632.7	14577.1	14925.8
COPA_HUMAN	Coatomer subunit alpha OS=Homo sapiens GN=COPA PE=1 SV=2	8	7	4.6	0.020	X	2757.3	3222.5	3151.5	181497.7	202001.2	236804.4	140673.0	138965.8	109032.7	52976.7	58477.9	60210.4	13108.4	19269.7	14315.0	1378.2	954.7	1724.3
SYQ_HUMAN	Glutamine--tRNA ligase OS=Homo sapiens GN=QARS PE=1 SV=1	2	2	4.5	0.022	X	16634.5	16552.9	15977.2	1497951.0	1409618.1	1335602.3	796202.1	849467.6	770251.2	443965.5	399966.2	380266.5	69258.8	86486.0	59414.1	12119.0	12243.7	12629.6
LARP1_HUMAN	La-related protein 1 OS=Homo sapiens GN=LARP1 PE=1 SV=2	4	2	4.3	0.037	X	75.9	44.4	12.9	15310.1	3771.1	11259.6	7222.4	8107.9	8707.2	2805.1	3856.4	4182.6	337.1	552.1	661.3	146.0	200.8	72.6
PDLI1_HUMAN	PDZ and LIM domain protein 1 OS=Homo sapiens GN=PDLIM1 PE=1 SV=4	3	2	4.2	0.031	X	2624.5	2635.6	2738.9	11443.8	16353.9	16889.8	2921.2	4256.0	3317.4	1327.5	1110.6	3217.9	1353.9	792.3	1103.3	2312.9	1886.3	1916.5
ACTN2_HUMAN	Alpha-actinin-2 OS=Homo sapiens GN=ACTN2 PE=1 SV=1	11	2	4.2	0.032	X	292.2	305.8	1404.6	159263.4	137404.9	157290.7	75344.0	78349.3	70425.7	43194.3	48230.6	47597.7	10450.9	10302.0	532.2	1197.4	411.7	361.4
DHX9_HUMAN	ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4	5	5	4.2	0.024	X	1058.9	1408.7	1353.3	98128.2	124838.8	112267.1	64552.3	77205.2	64144.7	37146.7	48539.6	33687.4	3378.6	3976.9	2830.5	529.5	425.9	557.0
ESYT1_HUMAN	Extended synaptotagmin-1 OS=Homo sapiens GN=ESYT1 PE=1 SV=1	6	5	4.1	0.036	X	2538.5	2813.4	2988.3	267541.3	321971.6	304635.0	122890.8	141963.9	148295.6	74801.6	103817.6	97336.0	10741.6	14847.1	12440.9	2364.3	2703.1	2188.3
LPPRC_HUMAN	Leucine-rich PPR motif-containing protein, mitochondrial OS=Homo sapiens GN=LRPPRC PE=1 SV=3	18	15	4.1	0.032	X	15084.5	15910.3	15349.4	1449756.8	1416403.8	1549205.2	680383.7	776543.2	714115.4	463247.3	484099.6	436372.8	48967.6	69089.6	71686.6	15177.6	15557.9	16438.5
HSP72_HUMAN	Heat shock-related 70 kDa protein 2 OS=Homo sapiens GN=HSPA2 PE=1 SV=1	53	2	4.1	0.027	X	615.3	520.3	605.7	29410.5	32992.7	38759.8	17697.2	19083.1	19432.6	11215.1	10970.5	11112.7	935.8	1954.5	1167.3	721.3	544.6	524.5
PSA_HUMAN	Puromycin-sensitive aminopeptidase OS=Homo sapiens GN=NPEPPS PE=1 SV=2	23	10	4.0	0.039	X	3888.9	4674.1	4630.3	474583.1	459713.9	464449.8	210245.4	226297.5	183882.9	163395.6	160987.4	146182.9	12247.1	6057.7	7635.1	2605.2	2609.8	2524.4
ITB1_HUMAN	Integrin beta-1 OS=Homo sapiens GN=ITGB1 PE=1 SV=2	7	5	4.0	0.038	X	1875.8	1832.3	2180.2	217691.7	201825.5	218186.2	88067.0	96543.8	92061.9	63382.5	63784.0	61779.8	12863.3	13425.8	8922.4	1390.9	1655.3	1415.7
CAPR1_HUMAN	Caprin-1 OS=Homo sapiens GN=CAPRIN1 PE=1 SV=2	5	5	4.0	0.033	X	6738.7	6944.9	6633.4	520627.4	603720.5	589372.3	249852.2	289076.3	294940.8	176545.1	170803.5	160083.4	34001.9	50404.4	35485.4	5018.6	4700.5	4549.7
LRC47_HUMAN	Leucine-rich repeat-containing protein 47 OS=Homo sapiens GN=LRRC47 PE=1 SV=1	2	1	4.0	0.047	X	362.6	340.1	387.2	42420.6	42244.7	40082.5	15369.7	15852.7	16917.7	14178.7	15004.8	11776.5	847.7	697.7	640.4	239.3	141.4	225.9
MSH2_HUMAN	DNA mismatch repair protein Msh2 OS=Homo sapiens GN=MSH2 PE=1 SV=1	3	1	4.0	0.036	X	912.3	866.6	821.6	98335.1	106554.2	117447.9	40402.0	49919.6	62893.3	29141.1	34230.1	28017.7	8528.3	9776.6	8710.3	820.7	837.8	670.2
TIF1B_HUMAN	Transcription intermediary factor 1-beta OS=Homo sapiens GN=TRIM28 PE=1 SV=5	21	15	3.9	0.036	X	15047.5	16442.0	16026.7	1802665.1	1886111.6	1826846.4	848049.6	894145.1	859703.6	579269.8	637778.2	570698.2	86772.9	93658.8	74371.6	10048.3	10053.8	10086.2
EIF3B_HUMAN	Eukaryotic translation initiation factor 3 subunit B OS=Homo sapiens GN=EIF3B PE=1 SV=3	7	5	3.9	0.027	X	4112.0	3902.7	3891.0	303657.4	304171.2	325887.3	178700.7	179616.2	190721.1	109403.0	119927.8	109174.9	13547.8	10007.9	5500.2	4561.3	4232.4	4035.9
VPS35_HUMAN	Vacuolar protein sorting-associated protein 35 OS=Homo sapiens GN=VPS35 PE=1 SV=2	4	4	3.9	0.026	X	123.9	150.7	339.3	44428.7	42423.1	61578.6	57660.4	61891.9	78106.1	31744.2	26285.8	28598.4	35.8	1056.1	232.5	73.9	196.5	132.8
UBA1_HUMAN	Ubiquitin-like modifier-activating enzyme 1 OS=Homo sapiens GN=UBA1 PE=1 SV=3	46	39	3.9	0.036	X	61951.9	60830.5	63359.7	6838083.0	6700813.0	6807245.7	3088893.1	3227652.4	3500060.4	2250489.2	2387795.4	2292083.7	227533.6	271084.7	255601.0	45959.0	45588.9	46787.0
DPP3_HUMAN	Dipeptidyl peptidase 3 OS=Homo sapiens GN=DPP3 PE=1 SV=2	9	6	3.8	0.027	X	709.4	746.8	734.8	86497.6	102262.2	98170.7	59382.2	68774.2	61125.9	40324.5	43327.3	34864.4	3030.3	687.5	2303.5	699.7	690.2	660.8
RENT1_HUMAN	Regulator of nonsense transcripts 1 OS=Homo sapiens GN=UPF1 PE=1 SV=2	10	7	3.8	0.026	X	5284.0	5416.2	5656.4	252057.3	263481.2	277501.4	145400.6	162794.8	161715.0	85797.3	102681.7	94914.3	14725.4	17556.2	12603.8	3790.7	3753.9	3858.3
NCAP_JUNIN	Nucleoprotein OS=Junin mammarenavirus GN=N PE=1 SV=1	2	1	3.8	0.044	X	1292.6	1419.5	1360.2	191230.8	223802.8	242300.8	85579.1	102005.2	105934.7	69399.5	81182.7	72825.1	8951.5	8708.2	7914.9	1478.0	1576.6	1520.5
HS90B_HUMAN	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4	87	24	3.8	0.037	X	56373.0	57669.8	59861.4	6826349.0	7403817.2	6741852.0	3384931.2	3416577.1	3671274.9	2626042.8	2382580.6	2505440.9	216562.0	275494.6	244970.1	53835.0	53456.6	55785.6
SFPQ_HUMAN	Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPQ PE=1 SV=2	13	12	3.8	0.035	X	7349.7	6956.3	7158.3	599303.7	619517.2	623334.5	272035.8	313636.1	338942.7	201778.9	215770.5	204957.1	36311.5	34210.6	31823.9	5823.8	5825.5	5566.7
ACTN1_HUMAN	Alpha-actinin-1 OS=Homo sapiens GN=ACTN1 PE=1 SV=2	16	6	3.7	0.021	X	1047.8	858.6	1160.6	37389.8	59108.7	54129.8	45335.9	40789.6	48623.3	16627.6	20972.0	25335.9	4165.1	2766.2	4451.8	1061.3	877.4	940.7
ENPL_HUMAN	Endoplasmin OS=Homo sapiens GN=HSP90B1 PE=1 SV=1	51	38	3.7	0.038	X	39290.7	40481.0	41708.0	4508992.4	4667041.2	4811426.9	2128427.8	2385800.6	2416587.3	1621455.8	1769377.6	1678656.0	151036.0	193227.8	134122.5	34917.2	33483.1	33238.2
KINH_HUMAN	Kinesin-1 heavy chain OS=Homo sapiens GN=KIF5B PE=1 SV=1	19	13	3.7	0.027	X	9985.2	10669.7	10406.5	561650.5	507780.7	479314.2	314863.7	354703.5	274729.2	159364.6	225283.3	187625.6	28205.3	29407.9	25188.6	8184.9	8289.6	8336.6
C1TC_HUMAN	C-1-tetrahydrofolate synthase, cytoplasmic OS=Homo sapiens GN=MTHFD1 PE=1 SV=3	36	30	3.7	0.033	X	16955.4	16511.7	17289.2	1528134.5	1403172.2	1485209.1	731638.1	797979.3	786700.8	486490.3	526692.4	466902.8	100127.8	115273.3	93197.9	14336.1	14467.0	14000.5
HNRL1_HUMAN	Heterogeneous nuclear ribonucleoprotein U-like protein 1 OS=Homo sapiens GN=HNRNPUL1 PE=1 SV=2	2	2	3.7	0.040	X	419.7	458.4	452.2	68566.0	46796.6	51489.7	28685.9	24494.3	32646.8	18879.3	25746.6	10422.2	4169.0	5524.2	2583.3	407.6	412.4	462.7
RBBP7_HUMAN	Histone-binding protein RBBP7 OS=Homo sapiens GN=RBBP7 PE=1 SV=1	4	2	3.7	0.045	X	3074.3	2849.1	2953.4	363347.3	405657.1	408781.2	164100.4	184083.1	177186.2	133230.8	142970.1	140758.8	13649.1	16846.3	12836.1	2448.0	2355.1	2442.5
MCM6_HUMAN	DNA replication licensing factor MCM6 OS=Homo sapiens GN=MCM6 PE=1 SV=1	10	8	3.6	0.042	X	2953.9	3071.2	2834.9	239532.5	275199.4	287703.4	102011.4	132145.9	128987.2	73679.3	90316.7	86520.5	21846.6	19653.8	20858.8	2895.2	3391.2	2723.9
PARP1_HUMAN	Poly [ADP-ribose] polymerase 1 OS=Homo sapiens GN=PARP1 PE=1 SV=4	32	27	3.6	0.031	X	21965.7	21126.3	22078.8	1399759.3	1355305.1	1414786.5	695611.2	827582.2	812385.0	479258.3	480081.7	474071.3	103043.2	97369.7	111017.7	20373.0	20120.4	20136.6
EZRI_HUMAN	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4	26	15	3.6	0.030	X	3779.3	3509.8	3135.8	361937.0	380186.4	412711.8	221219.3	243075.6	229519.9	133867.2	152731.7	141225.5	30339.1	21531.0	19604.4	4779.1	4654.5	4558.7
HS105_HUMAN	Heat shock protein 105 kDa OS=Homo sapiens GN=HSPH1 PE=1 SV=1	19	13	3.6	0.033	X	7132.5	7224.8	7549.0	640778.3	637577.0	616942.0	322196.5	347959.1	334534.3	202306.6	191608.5	182242.2	69777.6	77400.2	61884.6	7229.2	7430.5	7424.3
PSMD2_HUMAN	26S proteasome non-ATPase regulatory subunit 2 OS=Homo sapiens GN=PSMD2 PE=1 SV=3	8	7	3.6	0.034	X	3001.4	3568.6	3160.9	404033.3	423197.5	466768.7	229673.5	242505.5	256331.3	157628.5	177143.9	164615.6	17249.7	20086.6	15553.3	3119.7	3438.4	3169.4
TPIS_HUMAN	Triosephosphate isomerase OS=Homo sapiens GN=TPI1 PE=1 SV=3	21	21	3.6	0.026	X	2507.1	2229.9	2095.5	113988.9	126638.4	61653.9	72880.6	81601.4	81221.3	29566.8	30784.0	31639.0	18865.4	15913.9	15260.3	3107.9	3065.3	2763.2
CLH1_HUMAN	Clathrin heavy chain 1 OS=Homo sapiens GN=CLTC PE=1 SV=5	45	40	3.6	0.027	X	39140.5	39414.1	39445.8	2843422.8	2729297.0	2937101.6	1583705.2	1827467.2	1859566.0	886939.6	950675.1	874540.7	336412.7	374476.2	338371.7	35122.4	33468.5	34060.2
ACLY_HUMAN	ATP-citrate synthase OS=Homo sapiens GN=ACLY PE=1 SV=3	20	12	3.6	0.036	X	5840.9	6103.3	5531.0	566090.9	628783.4	609651.7	295027.6	331452.3	314577.1	201582.5	199643.7	188449.4	49160.7	66993.4	41996.3	7413.0	7878.1	7792.3
GANAB_HUMAN	Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB PE=1 SV=3	27	26	3.6	0.031	X	19628.0	19841.0	20467.8	1615476.3	1708817.8	1544191.4	940475.6	1035021.1	984101.9	647061.4	692666.3	640178.9	71347.2	62073.5	59403.0	14492.9	13628.9	13808.4
P5CS_HUMAN	Delta-1-pyrroline-5-carboxylate synthase OS=Homo sapiens GN=ALDH18A1 PE=1 SV=2	11	8	3.6	0.037	X	2205.5	2541.0	2525.4	355028.9	428523.9	481947.5	222463.4	294660.9	239839.5	153363.2	188506.5	197814.3	5578.8	7994.5	8678.4	2924.1	2674.0	3186.8
EIF3C_HUMAN	Eukaryotic translation initiation factor 3 subunit C OS=Homo sapiens GN=EIF3C PE=1 SV=1	9	3	3.5	0.045	X	465.2	588.9	519.8	59387.3	59943.3	59612.7	18848.7	29473.2	31378.4	14234.4	14844.1	17352.6	6167.9	9873.9	8360.6	956.3	1001.6	972.7
AT1A1_HUMAN	Sodium/potassium-transporting ATPase subunit alpha-1 OS=Homo sapiens GN=ATP1A1 PE=1 SV=1	19	6	3.5	0.034	X	6345.5	7137.0	7577.3	553534.2	596222.0	510331.7	320330.8	369100.4	295280.9	199540.7	244575.2	229905.8	23798.5	25662.4	22312.3	4615.9	4774.6	4783.0
CAND1_HUMAN	Cullin-associated NEDD8-dissociated protein 1 OS=Homo sapiens GN=CAND1 PE=1 SV=2	24	21	3.5	0.026	X	10945.4	12050.3	12618.5	1034656.4	1143221.5	1120168.2	730337.5	740156.9	788599.7	399023.0	411589.2	394582.0	105984.7	115881.7	125367.1	14222.1	14184.4	13708.1
DDB1_HUMAN	DNA damage-binding protein 1 OS=Homo sapiens GN=DDB1 PE=1 SV=1	5	5	3.5	0.028	X	3308.3	2850.1	3478.3	233582.0	231437.8	227483.8	122865.5	198355.0	159116.1	94083.0	86310.2	80108.7	39651.7	13519.9	17485.4	2738.2	2881.2	2584.8
CPSF6_HUMAN	Cleavage and polyadenylation specificity factor subunit 6 OS=Homo sapiens GN=CPSF6 PE=1 SV=2	6	6	3.5	0.021	X	1988.4	2021.1	2138.7	118082.4	110712.6	104052.3	110733.3	117177.8	116519.3	52208.5	49997.0	48872.6	13507.6	15934.8	10890.6	1651.2	1863.8	1614.6
CSDE1_HUMAN	Cold shock domain-containing protein E1 OS=Homo sapiens GN=CSDE1 PE=1 SV=2	5	2	3.5	0.033	X	842.1	1054.2	868.0	67273.2	81284.1	84225.1	45018.1	41404.5	48865.6	24171.0	26298.5	27582.8	8895.4	9088.9	7912.1	918.9	1053.7	865.7
IPO7_HUMAN	Importin-7 OS=Homo sapiens GN=IPO7 PE=1 SV=1	3	3	3.5	0.027	X	1586.2	1711.6	1924.8	92144.4	92691.6	100069.3	67330.0	67610.9	74870.3	43531.0	47572.2	41654.4	2899.9	3371.3	4059.3	661.1	514.2	593.3
PSD12_HUMAN	26S proteasome non-ATPase regulatory subunit 12 OS=Homo sapiens GN=PSMD12 PE=1 SV=3	5	3	3.5	0.020	X	1659.3	1650.9	1583.8	50162.5	61816.5	64844.2	51840.2	62283.6	66695.1	18153.5	23987.7	22624.7	11197.3	14570.5	11788.5	937.5	947.7	842.7
ACON_HUMAN	Aconitate hydratase, mitochondrial OS=Homo sapiens GN=ACO2 PE=1 SV=2	3	2	3.4	0.048	X	883.9	959.0	759.8	46536.1	59131.0	56821.8	25814.7	21705.8	26755.9	26501.1	17657.7	17250.1	1891.5	3598.3	1394.7	348.4	460.3	488.6
ACTN4_HUMAN	Alpha-actinin-4 OS=Homo sapiens GN=ACTN4 PE=1 SV=2	35	23	3.4	0.032	X	17431.8	17841.2	18031.1	1358850.6	1417543.9	1397197.7	823216.8	921778.4	888667.5	545342.6	612568.1	582698.7	70909.0	71249.5	83700.7	12910.7	12413.3	12934.0
EF2_HUMAN	Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4	93	86	3.4	0.040	X	131143.3	136435.6	133815.6	15864687.0	14843083.2	14695201.8	7384807.2	8514863.4	7917333.1	5450152.2	5972253.6	5394610.7	925071.6	1188448.0	1015253.8	173232.9	172971.7	174702.3
HNRPU_HUMAN	Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU PE=1 SV=6	22	17	3.4	0.045	X	27628.9	32188.8	30870.4	2856274.0	2912281.2	2945305.2	1277396.4	1445131.8	1468405.2	1012194.6	1173531.4	1058121.4	146058.8	182868.9	175700.4	26805.7	24969.8	26518.4
COPB2_HUMAN	Coatomer subunit beta' OS=Homo sapiens GN=COPB2 PE=1 SV=2	9	6	3.4	0.023	X	33084.8	33609.5	33464.9	1815736.9	1884715.0	1838145.6	1293518.3	1495625.0	1482098.2	562000.4	744822.9	634836.0	260332.4	325861.2	296174.6	31872.5	30591.2	32267.2
ADH1_YEAST	Alcohol dehydrogenase 1 OS=Saccharomyces cerevisiae (strain ATCC 204508 / S288c) GN=ADH1 PE=1 SV=5	59	55	3.4	0.048	X	207754.8	214458.9	223322.2	24865716.8	24770529.0	24896810.7	10135387.4	11456578.0	11282084.9	7465503.6	8253888.0	7322218.9	2416991.3	3073035.6	2685166.1	247110.7	239895.6	241580.4
TERA_HUMAN	Transitional endoplasmic reticulum ATPase OS=Homo sapiens GN=VCP PE=1 SV=4	34	29	3.4	0.044	X	23687.0	24433.6	24985.8	3238577.2	2922349.1	2838745.8	1402909.3	1610880.5	1516296.4	1197162.0	1172705.3	1126175.8	153637.2	152740.7	142366.7	26076.6	26709.6	26502.0
HXK1_HUMAN	Hexokinase-1 OS=Homo sapiens GN=HK1 PE=1 SV=3	6	5	3.4	0.026	X	8760.2	7907.5	8604.6	447534.9	430620.0	361132.5	295712.3	286421.5	297975.4	155627.6	162229.5	137733.0	49422.8	56314.9	48408.3	8596.1	9366.4	9266.0
AINX_HUMAN	Alpha-internexin OS=Homo sapiens GN=INA PE=1 SV=2	3	2	3.4	0.044	X	720.5	512.2	437.7	105759.1	128211.7	114490.0	63085.3	61491.9	48469.4	32454.6	35369.2	30203.2	18823.3	18786.5	15802.4	1517.8	1325.4	1378.4
XRCC5_HUMAN	X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3	20	17	3.4	0.033	X	5132.4	5130.6	4790.4	848938.7	855427.6	803484.3	541110.0	611379.1	621179.5	373247.9	409185.4	409762.5	23682.0	24197.3	22900.5	5496.8	5293.4	5288.4
STRAP_HUMAN	Serine-threonine kinase receptor-associated protein OS=Homo sapiens GN=STRAP PE=1 SV=1	4	3	3.3	0.047	X	1314.6	1180.6	1081.1	68957.5	61294.3	64777.9	24485.2	31368.0	28931.6	15757.0	18376.0	21220.8	7601.7	8100.0	7139.7	2170.6	2292.5	2121.8
TNPO1_HUMAN	Transportin-1 OS=Homo sapiens GN=TNPO1 PE=1 SV=2	8	7	3.3	0.043	X	1814.6	1813.9	1973.8	240599.8	274952.3	295713.5	130588.0	173274.1	151535.0	112393.1	126589.6	106074.9	11050.6	8036.4	10410.6	2433.5	1900.1	2058.2
SAE2_HUMAN	SUMO-activating enzyme subunit 2 OS=Homo sapiens GN=UBA2 PE=1 SV=2	4	2	3.3	0.036	X	292.8	353.5	360.4	88941.2	79459.2	83935.1	49749.0	59425.4	62865.2	36234.2	44044.5	40312.4	2764.6	1933.8	1767.7	277.8	323.2	313.9
MCM7_HUMAN	DNA replication licensing factor MCM7 OS=Homo sapiens GN=MCM7 PE=1 SV=4	12	10	3.3	0.030	X	1934.8	1714.6	2235.8	184622.3	211408.8	243250.3	127942.5	164956.6	185550.5	69603.7	86451.7	100367.6	20110.2	27192.9	25330.5	3356.9	3005.6	3168.9
SND1_HUMAN	Staphylococcal nuclease domain-containing protein 1 OS=Homo sapiens GN=SND1 PE=1 SV=1	17	15	3.3	0.042	X	7630.2	8230.7	7655.3	1264274.2	1069724.6	1178879.1	616193.7	650399.3	658757.1	495366.8	466424.4	421042.5	78042.4	86570.0	78043.9	7620.3	7892.0	7056.3
VINC_HUMAN	Vinculin OS=Homo sapiens GN=VCL PE=1 SV=4	40	34	3.3	0.034	X	31082.3	32866.3	32602.0	1839665.1	2013807.0	1839512.0	1108425.0	1181534.6	1127508.1	704916.5	753723.3	696308.8	164049.1	215412.4	171006.7	28614.2	28447.3	28308.6
GLRX3_HUMAN	Glutaredoxin-3 OS=Homo sapiens GN=GLRX3 PE=1 SV=2	11	9	3.3	0.028	X	3741.9	3690.0	3774.6	45872.3	24091.6	66745.5	69439.9	48943.7	48142.2	31945.8	10076.5	16547.0	7103.4	11312.2	7387.6	4055.6	3477.2	4342.3
HS90A_HUMAN	Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5	92	36	3.3	0.037	X	140960.0	143551.3	146854.8	7717781.1	7715400.6	8063598.0	4377563.4	4708810.3	4769639.2	3209068.3	3621536.7	3319869.6	295178.5	384821.0	323362.0	138739.1	137827.6	139505.8
LAP2B_HUMAN	Lamina-associated polypeptide 2, isoforms beta/gamma OS=Homo sapiens GN=TMPO PE=1 SV=2	8	2	3.3	0.042	X	49.9	30.3	189.2	5968.1	4900.2	3808.8	6457.0	2853.1	7293.2	3712.9	2105.9	2515.5	183.2	439.5	167.7	122.7	229.1	226.5
MYH9_HUMAN	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4	63	39	3.2	0.028	X	27508.1	28528.4	28619.0	1741109.0	1596378.2	1648948.9	1273273.1	1310063.3	1204847.6	702490.7	742791.6	697781.8	171720.4	201705.3	151733.1	20247.4	19878.6	19384.7
MYH14_HUMAN	Myosin-14 OS=Homo sapiens GN=MYH14 PE=1 SV=2	16	10	3.2	0.044	X	10276.2	10870.7	10707.2	1266542.6	1317935.8	1322433.6	565470.3	926712.4	649567.3	394921.7	654092.4	412618.7	98060.7	120406.6	169691.1	10079.8	10248.4	10153.6
MCM5_HUMAN	DNA replication licensing factor MCM5 OS=Homo sapiens GN=MCM5 PE=1 SV=5	11	9	3.2	0.036	X	2607.7	2570.9	2244.8	316535.1	292178.5	292180.3	200752.8	243979.2	234046.1	140121.9	166665.7	167831.7	4254.8	5649.3	3323.3	2553.1	2713.4	2769.6
VDAC1_HUMAN	Voltage-dependent anion-selective channel protein 1 OS=Homo sapiens GN=VDAC1 PE=1 SV=2	10	7	3.2	0.029	X	8109.9	7764.7	7763.3	191718.1	198648.6	193109.4	114875.9	119762.3	132874.8	76660.9	81647.7	86203.2	9118.7	10760.1	10798.2	9805.7	9916.3	10197.7
ACOT9_HUMAN	Acyl-coenzyme A thioesterase 9, mitochondrial OS=Homo sapiens GN=ACOT9 PE=1 SV=2	7	6	3.2	0.026	X	2742.7	3443.8	2350.6	75696.2	117089.5	126053.9	99933.1	103046.3	106717.7	27360.4	32892.8	28771.3	32206.6	36890.6	33915.9	2984.2	3148.2	2873.3
SF01_HUMAN	Splicing factor 1 OS=Homo sapiens GN=SF1 PE=1 SV=4	7	6	3.1	0.028	X	3110.7	3788.1	2785.2	147511.5	194060.4	211495.6	158039.7	196937.6	190975.6	65978.5	98575.1	92536.8	27197.0	29367.7	27331.5	4315.1	3569.7	3322.0
DHX15_HUMAN	Pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15 OS=Homo sapiens GN=DHX15 PE=1 SV=2	6	4	3.1	0.041	X	1993.8	2182.4	2219.6	226230.5	248443.3	210392.7	145909.8	151160.4	150355.3	115168.3	113056.5	105128.1	7240.3	8037.8	7866.4	2187.3	1908.5	2136.9
STX11_HUMAN	Syntaxin-11 OS=Homo sapiens GN=STX11 PE=1 SV=1	4	4	3.1	0.029	X	883.5	1053.0	945.4	47194.4	40461.5	51956.3	38584.1	36974.6	36450.8	18529.5	24514.3	18625.4	5750.1	6849.9	4257.0	842.0	885.8	869.2
MIC60_HUMAN	MICOS complex subunit MIC60 OS=Homo sapiens GN=IMMT PE=1 SV=1	8	5	3.1	0.038	X	2536.2	2711.8	2627.6	236837.7	233677.5	247945.4	151179.1	180873.2	171752.4	119503.3	129095.1	109061.6	6887.8	10721.3	9829.9	2949.7	2503.2	2557.4
IPYR_HUMAN	Inorganic pyrophosphatase OS=Homo sapiens GN=PPA1 PE=1 SV=2	6	6	3.1	0.039	X	3548.0	4088.3	3957.8	185637.5	203393.8	203791.2	116838.2	108405.9	102870.0	66048.0	73965.9	71062.5	24203.8	26888.6	20472.8	5910.5	4609.3	5090.7
DHE3_HUMAN	Glutamate dehydrogenase 1, mitochondrial OS=Homo sapiens GN=GLUD1 PE=1 SV=2	12	6	3.1	0.032	X	4244.5	4023.5	3348.9	303131.9	259882.1	262214.9	195198.4	306371.6	207625.8	125936.1	137179.0	115971.5	29624.3	34819.0	41941.8	3347.6	3124.0	3100.4
COPG1_HUMAN	Coatomer subunit gamma-1 OS=Homo sapiens GN=COPG1 PE=1 SV=1	8	6	3.1	0.044	X	2045.0	2076.8	2060.9	213917.1	202917.7	193638.5	119749.7	194292.2	114656.5	65616.5	150501.5	75467.5	12793.5	9954.6	16453.2	2028.0	2091.2	1917.0
U5S1_HUMAN	116 kDa U5 small nuclear ribonucleoprotein component OS=Homo sapiens GN=EFTUD2 PE=1 SV=1	18	6	3.1	0.043	X	3387.8	3367.0	3418.3	128155.8	128918.1	138515.0	61712.5	91705.6	128816.3	51505.1	94577.5	55772.8	4399.6	7523.2	3574.8	2221.6	2146.5	2366.5
PDIP2_HUMAN	Polymerase delta-interacting protein 2 OS=Homo sapiens GN=POLDIP2 PE=1 SV=1	2	1	3.1	0.029	X	2028.4	2329.5	2360.5	54194.8	52150.6	57366.0	58357.0	64774.4	56695.7	32130.4	36904.1	33425.7	3208.4	2275.1	3109.5	1191.6	1234.1	1266.3
EF1B_HUMAN	Elongation factor 1-beta OS=Homo sapiens GN=EEF1B2 PE=1 SV=3	2	2	3.1	0.045	X	587.9	653.1	592.0	24645.3	21482.4	23611.7	25412.5	23062.1	7461.7	8430.0	11495.2	14579.8	826.5	502.0	452.8	2029.9	1636.9	1847.2
NUCL_HUMAN	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3	43	40	3.0	0.043	X	29815.6	31088.4	30808.3	3815564.7	3913156.0	3516674.9	2312512.8	2479183.0	2336456.9	1656108.8	1883489.3	1765787.9	175772.9	238876.5	183804.0	55221.5	52964.2	53637.1
AIP_HUMAN	AH receptor-interacting protein OS=Homo sapiens GN=AIP PE=1 SV=2	3	3	3.0	0.040	X	3100.0	3293.4	3223.0	136974.2	182175.1	179220.6	91306.2	100817.4	112428.1	55311.1	66272.9	61041.0	21819.2	27456.5	25252.5	4056.5	4268.6	3863.7
DDX17_HUMAN	Probable ATP-dependent RNA helicase DDX17 OS=Homo sapiens GN=DDX17 PE=1 SV=2	29	17	3.0	0.040	X	12514.2	12816.6	12937.8	629531.4	658856.2	718699.9	420463.7	449204.8	415447.0	288344.4	345834.9	295174.6	37402.0	46950.2	51131.0	14485.8	13716.8	13951.8
RIR1_HUMAN	Ribonucleoside-diphosphate reductase large subunit OS=Homo sapiens GN=RRM1 PE=1 SV=1	10	8	3.0	0.037	X	1367.2	1224.1	1374.5	218836.3	209139.8	221138.7	215516.5	194060.3	204683.1	107298.7	151932.3	118937.3	9556.0	14098.3	13998.8	2387.3	2591.1	2245.0
FUS_HUMAN	RNA-binding protein FUS OS=Homo sapiens GN=FUS PE=1 SV=1	5	3	3.0	0.047	X	12597.9	12835.6	12708.3	455500.1	429371.5	426643.8	228139.6	234035.8	237988.5	182798.1	212309.7	180548.0	28977.3	29815.7	30052.2	9507.8	10170.4	9665.3
PRDX1_HUMAN	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1	4	2	2.9	0.040	X	442.6	464.1	513.1	20959.7	27710.1	27422.9	15656.4	17326.9	28627.7	4994.9	13230.5	8955.2	4561.1	8420.0	5284.8	988.4	827.5	632.0
UBQL1_HUMAN	Ubiquilin-1 OS=Homo sapiens GN=UBQLN1 PE=1 SV=2	2	1	2.9	0.046	X	611.4	679.6	443.1	81921.2	71157.2	68301.6	43940.4	50092.5	47474.4	31234.9	37531.7	27735.2	8343.6	9527.3	10128.4	854.1	682.2	567.2
NSUN2_HUMAN	tRNA (cytosine(34)-C(5))-methyltransferase OS=Homo sapiens GN=NSUN2 PE=1 SV=2	9	6	2.9	0.033	X	2085.5	2130.3	2202.0	86434.0	92639.8	101827.2	106168.6	110845.5	107334.6	45245.4	56659.9	45888.1	20732.6	19412.6	19334.7	2035.4	1858.8	2110.8
FUBP1_HUMAN	Far upstream element-binding protein 1 OS=Homo sapiens GN=FUBP1 PE=1 SV=3	13	9	2.9	0.035	X	8575.3	8269.2	8380.8	289457.3	281780.6	285356.0	211567.7	198802.7	192288.2	121639.1	139041.2	111293.1	29459.2	56488.8	24789.9	11879.5	12962.9	12574.7
IMB1_HUMAN	Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2	25	23	2.9	0.044	X	69779.7	85022.4	93288.2	5927141.1	8075634.6	7874792.7	3964839.7	4778440.7	5916608.0	2290993.8	2463422.0	2983673.0	1317163.0	1560194.3	1827470.9	145501.3	155006.0	152701.9
PHB2_HUMAN	Prohibitin-2 OS=Homo sapiens GN=PHB2 PE=1 SV=2	18	17	2.8	0.032	X	17945.1	18805.6	18048.5	203454.1	109480.4	248935.8	138023.4	118255.4	127695.1	58920.0	85391.1	56103.0	25019.3	33379.2	32610.5	19325.3	21022.7	20177.3
ACTZ_HUMAN	Alpha-centractin OS=Homo sapiens GN=ACTR1A PE=1 SV=1	5	4	2.8	0.012	X	1899.2	1812.0	1902.7	13177.7	8932.3	11060.5	9708.2	12886.2	9583.8	1970.1	1843.0	3227.1	4811.3	2658.5	5402.4	1752.3	1844.7	1802.6
ROA1_HUMAN	Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 PE=1 SV=5	7	4	2.8	0.046	X	10561.7	10643.3	10764.0	31227.5	37325.1	45007.0	4916.2	11420.3	5369.3	5633.9	3019.2	9059.6	6161.9	2956.9	3132.6	9845.9	10868.5	9561.7
EIF3A_HUMAN	Eukaryotic translation initiation factor 3 subunit A OS=Homo sapiens GN=EIF3A PE=1 SV=1	24	16	2.7	0.044	X	7313.8	8257.8	7388.3	332269.5	343526.3	312530.1	284140.8	337775.9	402314.3	176703.4	246006.3	201532.1	33818.1	37727.7	35010.1	3873.3	4153.0	3963.3
STAU1_HUMAN	Double-stranded RNA-binding protein Staufen homolog 1 OS=Homo sapiens GN=STAU1 PE=1 SV=2	4	3	2.7	0.049	X	1541.9	1715.3	1698.9	125281.4	136267.5	109001.6	91467.6	89700.4	95166.0	59300.6	74133.0	65812.9	11282.3	9073.8	12317.9	2231.8	2201.4	2228.5
PYGL_HUMAN	Glycogen phosphorylase, liver form OS=Homo sapiens GN=PYGL PE=1 SV=4	8	4	2.7	0.044	X	2713.9	2684.5	2218.1	84750.2	105744.9	72324.0	68341.1	75442.2	92641.6	44592.0	55671.8	49186.3	5470.1	12724.1	12804.9	2205.9	1894.9	1789.0
FETUA_HUMAN	Alpha-2-HS-glycoprotein OS=Homo sapiens GN=AHSG PE=1 SV=1	8	7	2.7	0.043	X	12576.7	12808.7	11807.5	591715.2	591015.2	599922.0	377928.0	493202.3	415146.8	243080.5	315447.8	247707.1	95364.4	109766.6	104467.2	11377.5	12320.5	10483.7
SEPT9_HUMAN	Septin-9 OS=Homo sapiens GN=SEPT9 PE=1 SV=2	5	3	2.7	0.037	X	9417.6	9348.1	9671.8	117233.8	74384.1	128735.4	98742.1	62228.4	62907.9	37867.2	56166.5	50014.8	12035.1	15576.0	9625.5	10438.6	10609.3	10697.3
G3PT_HUMAN	Glyceraldehyde-3-phosphate dehydrogenase, testis-specific OS=Homo sapiens GN=GAPDHS PE=1 SV=2	2	1	2.7	0.000	X	19314.1	19885.3	20181.3	34799.5	32124.3	33892.7	17243.2	14410.4	22078.0	14344.2	10573.4	8327.0	6575.9	5778.4	4681.3	9680.2	9594.5	10290.2
HSP74_HUMAN	Heat shock 70 kDa protein 4 OS=Homo sapiens GN=HSPA4 PE=1 SV=4	22	17	2.7	0.044	X	44628.1	44024.8	44003.5	909580.4	920157.6	793023.3	502805.6	501703.2	516501.5	360918.0	367345.2	375801.5	89051.9	91739.7	98338.3	71874.5	72241.9	72938.1
PCBP1_HUMAN	Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2	11	7	2.7	0.035	X	7329.8	7343.6	7814.9	94101.0	101051.2	109199.6	56662.8	62450.7	60956.6	33146.1	37857.2	33902.6	19937.7	25234.5	21124.8	6671.1	6604.2	5929.6
GDIA_HUMAN	Rab GDP dissociation inhibitor alpha OS=Homo sapiens GN=GDI1 PE=1 SV=2	11	2	2.6	0.043	X	762.2	841.2	923.2	7723.8	5115.5	4313.4	9130.1	11147.5	11415.5	4865.9	3443.0	2601.3	1430.3	420.6	1175.5	1803.2	2056.4	1717.3
SRSF7_HUMAN	Serine/arginine-rich splicing factor 7 OS=Homo sapiens GN=SRSF7 PE=1 SV=1	2	2	2.6	0.030	X	2626.0	2790.4	2565.8	16348.8	27505.1	27357.9	16971.1	26797.6	22887.7	6868.3	15305.2	13007.2	4599.1	6959.6	4088.8	1930.8	1517.6	1506.7
RL8_HUMAN	60S ribosomal protein L8 OS=Homo sapiens GN=RPL8 PE=1 SV=2	15	12	2.6	0.048	X	399.2	377.8	282.0	9497.4	13798.3	14923.5	7320.8	8645.4	12048.1	5431.3	5010.5	2824.8	2283.8	2390.4	2958.0	1703.0	1884.5	1443.7
BACH_HUMAN	Cytosolic acyl coenzyme A thioester hydrolase OS=Homo sapiens GN=ACOT7 PE=1 SV=3	4	4	2.6	0.035	X	2335.4	2409.4	2672.6	17099.9	23343.7	25242.5	11960.0	14007.9	17990.7	11038.1	9573.1	13437.9	2457.3	1910.7	3324.0	1084.5	1135.4	1150.7
ANXA2_HUMAN	Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2	25	6	2.6	0.040	X	6005.7	5773.4	6057.7	161450.0	147169.2	166310.4	136765.3	125022.5	106179.7	71661.8	53935.5	60343.9	36635.4	56769.0	43936.0	3834.8	4019.9	3776.8
HS74L_HUMAN	Heat shock 70 kDa protein 4L OS=Homo sapiens GN=HSPA4L PE=1 SV=3	10	8	2.5	0.046	X	3821.7	4746.3	4734.3	108884.8	116856.2	108711.8	92665.4	132162.0	76090.2	46200.7	75813.2	55596.1	14119.5	18128.1	21772.8	8145.0	7237.4	7661.3
LDHC_HUMAN	L-lactate dehydrogenase C chain OS=Homo sapiens GN=LDHC PE=1 SV=4	3	1	2.5	0.044	X	6774.5	6475.3	6956.0	404391.2	444000.1	449593.5	420779.2	455167.7	477557.8	220909.3	240623.9	219278.2	104077.1	115948.1	107553.8	13864.3	13945.4	13847.8
PRS10_HUMAN	26S protease regulatory subunit 10B OS=Homo sapiens GN=PSMC6 PE=1 SV=1	3	3	2.5	0.027	X	1472.0	1433.6	1480.9	9986.0	11196.8	12849.2	5514.7	10713.4	11553.8	2032.4	4812.4	6074.4	3150.8	3131.0	3401.8	1110.1	1174.7	1184.0
KCRB_HUMAN	Creatine kinase B-type OS=Homo sapiens GN=CKB PE=1 SV=1	15	13	2.5	0.035	X	16864.8	16371.9	17104.9	205986.9	239744.4	288145.6	218747.2	211528.0	202666.7	118450.0	125474.5	117347.1	50824.6	55674.7	50577.7	15768.2	15733.7	15996.2
TBB2A_HUMAN	Tubulin beta-2A chain OS=Homo sapiens GN=TUBB2A PE=1 SV=1	55	2	2.5	0.040	X	4639.9	4656.8	4800.5	120700.0	123595.6	110564.0	99804.5	123505.0	115896.5	62810.3	65920.4	51873.9	27286.4	28409.1	30436.3	4761.6	5593.7	5821.7
LKHA4_HUMAN	Leukotriene A-4 hydrolase OS=Homo sapiens GN=LTA4H PE=1 SV=2	3	2	2.5	0.048	X	3324.2	3666.2	4019.7	360988.5	342553.0	354658.6	275429.7	370369.6	362001.9	141622.5	176578.9	160895.0	75998.1	141991.1	99652.1	12440.0	11110.9	12758.4
CHM4B_HUMAN	Charged multivesicular body protein 4b OS=Homo sapiens GN=CHMP4B PE=1 SV=1	2	2	2.5	0.043	X	1069.1	1333.3	987.2	7960.1	9316.2	5348.2	12333.8	8195.1	13029.0	6815.2	5308.8	3141.4	1730.5	2037.4	2048.5	1037.9	1137.8	1027.3
MCMBP_HUMAN	Mini-chromosome maintenance complex-binding protein OS=Homo sapiens GN=MCMBP PE=1 SV=2	1	1	22.5	0.143		36.7	51.0	38.6	9045.6	4887.6	26802.0	979.3	1329.1	1301.5	20.6	1206.3	65.2	69.0	305.9	246.8	13.6	25.0	27.0
GSTO1_HUMAN	Glutathione S-transferase omega-1 OS=Homo sapiens GN=GSTO1 PE=1 SV=2	2	1	18.3	0.264		0.2	0.0	0.0	0.0	533.6	0.0	0.0	0.0	146.7	34.5	0.0	0.0	0.0	0.0	0.0	0.0	2.1	0.5
RADI_HUMAN	Radixin OS=Homo sapiens GN=RDX PE=1 SV=1	16	3	12.5	0.280		3705.6	4133.2	3831.1	36847.8	817745.3	22908.8	35856.7	39183.2	38571.2	14003.0	16531.9	17775.0	3209.3	15660.2	2248.1	2960.0	3682.4	3965.3
SNX5_HUMAN	Sorting nexin-5 OS=Homo sapiens GN=SNX5 PE=1 SV=1	2	2	12.2	0.252		888.3	906.3	952.5	4676.2	77145.1	3533.4	4245.3	3725.5	3964.3	1540.7	668.0	536.3	2095.1	1433.6	1240.6	63.3	279.4	359.2
T2FA_HUMAN	General transcription factor IIF subunit 1 OS=Homo sapiens GN=GTF2F1 PE=1 SV=2	2	2	11.6	0.066		4977.5	5808.7	6440.8	127600.4	113525.1	99808.7	10886.3	2751.3	6724.2	10096.9	8043.2	1479.8	766.3	569.6	629.4	2922.9	3963.9	4159.4
FHL1_HUMAN	Four and a half LIM domains protein 1 OS=Homo sapiens GN=FHL1 PE=1 SV=4	1	1	10.3	0.022		128.4	94.8	88.3	40123.0	20692.9	51897.9	14662.4	59438.7	66526.1	9531.7	7725.8	1177.7	1201.1	392.4	3919.5	248.8	237.3	225.3
GEMI_HUMAN	Geminin OS=Homo sapiens GN=GMNN PE=1 SV=1	1	1	7.5	0.235		48.2	25.0	25.8	0.0	0.0	0.0	0.0	0.0	219.6	0.0	0.0	0.0	21.5	0.0	0.0	21.1	0.0	0.0
VIGLN_HUMAN	Vigilin OS=Homo sapiens GN=HDLBP PE=1 SV=2	4	3	7.1	0.052		297.1	317.5	284.2	63671.1	39865.0	91722.8	16580.6	20393.1	20549.9	14016.6	9157.5	11851.9	51.9	435.6	266.0	2.3	10.3	0.0
H90B3_HUMAN	Putative heat shock protein HSP 90-beta-3 OS=Homo sapiens GN=HSP90AB3P PE=5 SV=1	35	2	7.0	0.092		329.0	334.5	238.2	1915.0	1016.6	2741.9	26799.2	29148.2	24778.1	3600.9	4078.7	1646.4	1243.0	607.1	310.9	394.5	302.4	344.7
H2BFM_HUMAN	Histone H2B type F-M OS=Homo sapiens GN=H2BFM PE=1 SV=2	1	1	6.9	0.188		15.3	21.1	4.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	5.9	0.0
1433G_HUMAN	14-3-3 protein gamma OS=Homo sapiens GN=YWHAG PE=1 SV=2	4	1	6.4	0.425		0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	514.5	0.0	0.0	0.0	0.0	0.0	57.1	16.0	0.0	7.0
TBA1A_HUMAN	Tubulin alpha-1A chain OS=Homo sapiens GN=TUBA1A PE=1 SV=1	56	2	5.2	0.063		55.2	38.2	32.8	18967.2	20940.0	16252.7	3343.5	3546.7	6398.6	3217.5	2657.7	5867.8	500.4	276.1	550.1	90.2	117.8	87.3
IF4B_HUMAN	Eukaryotic translation initiation factor 4B OS=Homo sapiens GN=EIF4B PE=1 SV=2	3	2	5.2	0.110		0.0	194.1	170.5	28115.9	21611.9	20871.1	3094.4	1857.6	986.5	1327.0	5108.2	3681.0	1824.5	217.9	2053.6	192.9	267.4	146.5
HNRPM_HUMAN	Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3	1	1	4.7	0.048		285.9	355.8	251.6	23698.7	31972.2	34775.0	7942.6	12546.4	9420.3	5356.3	8476.0	4927.6	1681.4	1890.5	2401.5	366.5	345.7	184.1
SC24C_HUMAN	Protein transport protein Sec24C OS=Homo sapiens GN=SEC24C PE=1 SV=3	5	3	4.7	0.070		603.9	730.7	519.0	225058.3	217512.8	156979.7	50508.9	47819.3	39231.5	32649.5	39570.0	27409.1	15923.8	16736.2	15998.5	2719.9	2643.5	2803.7
LDH6B_HUMAN	L-lactate dehydrogenase A-like 6B OS=Homo sapiens GN=LDHAL6B PE=1 SV=3	2	1	4.5	0.124		4.2	0.0	6.2	812.1	0.0	652.4	12.2	294.6	302.5	0.0	105.1	37.7	147.8	90.7	37.6	18.1	10.8	11.5
ACTBL_HUMAN	Beta-actin-like protein 2 OS=Homo sapiens GN=ACTBL2 PE=1 SV=2	26	1	4.3	0.051		3753.1	4059.4	3396.8	722914.2	751715.6	779298.1	234845.2	245221.8	259106.8	177760.4	198475.4	165294.0	49980.4	51817.3	43783.7	5887.3	5137.0	5221.7

XRN2_HUMAN	5'-3' exoribonuclease 2 OS=Homo sapiens GN=XRN2 PE=1 SV=1	1	1	4.2	0.030		6.5	18.3	0.0	11258.8	6359.1	17342.2	9596.4	13061.3	8781.6	5177.4	3526.6	5176.0	923.6	898.3	159.8	30.2	0.0	18.7
RL7A_HUMAN	60S ribosomal protein L7a OS=Homo sapiens GN=RPL7A PE=1 SV=2	10	8	4.1	0.050		516.3	456.0	778.2	32758.4	21767.3	61294.7	22736.4	25499.5	16566.3	5898.6	6050.9	22431.5	2612.4	3121.2	1960.5	624.7	780.8	666.7
IF4G1_HUMAN	Eukaryotic translation initiation factor 4 gamma 1 OS=Homo sapiens GN=EIF4G1 PE=1 SV=4	13	11	4.0	0.063		6929.1	7182.2	6794.2	491481.7	654136.1	789627.5	196242.4	188897.8	232072.5	148423.8	172825.5	188052.1	37530.0	43943.3	40852.4	5643.2	5608.9	5314.2
DCTN2_HUMAN	Dynactin subunit 2 OS=Homo sapiens GN=DCTN2 PE=1 SV=4	7	7	4.0	0.117		1408.7	1498.0	1341.2	379.3	1552.6	8078.3	33987.4	40792.6	32747.3	887.7	343.1	321.6	8485.4	9709.1	6991.0	1506.2	1100.2	1403.1
PGTB1_HUMAN	Geranylgeranyl transferase type-1 subunit beta OS=Homo sapiens GN=PGGT1B PE=1 SV=2	4	3	4.0	0.075		2926.8	3655.0	3720.1	508960.3	542557.2	580505.5	116916.1	146354.1	138296.7	137234.1	139597.8	124459.2	40052.2	37809.4	32063.0	1809.5	2037.5	1698.6
CDK1_HUMAN	Cyclin-dependent kinase 1 OS=Homo sapiens GN=CDK1 PE=1 SV=3	5	5	3.9	0.052		4816.9	4237.0	4078.9	879701.0	863168.6	880260.5	305146.9	308956.3	285950.0	195267.8	199957.4	205853.2	83504.0	91304.9	83529.8	12835.6	12596.2	13328.3
ACTN3_HUMAN	Alpha-actinin-3 OS=Homo sapiens GN=ACTN3 PE=1 SV=2	9	1	3.9	0.140		202.3	251.1	2484.2	9027.1	1038.2	1026.9	53007.1	64174.7	61975.9	13361.7	13636.8	11295.5	2213.3	2830.3	2840.2	1063.0	1063.4	1117.6
PGAM1_HUMAN	Phosphoglycerate mutase 1 OS=Homo sapiens GN=PGAM1 PE=1 SV=2	9	3	3.9	0.133		199.6	168.8	101.3	19882.6	11437.3	43603.1	5563.7	12124.4	3817.4	4955.7	11526.9	4635.4	936.0	1480.3	1239.4	63.5	82.5	88.9
PFKAP_HUMAN	ATP-dependent 6-phosphofructokinase, platelet type OS=Homo sapiens GN=PFKP PE=1 SV=2	3	3	3.7	0.053		1790.8	1914.5	2352.7	294620.2	343180.5	351601.7	87799.0	139881.2	145129.8	12626.2	81534.7	85856.7	51628.2	70854.0	54715.1	3661.5	3763.5	3177.3
E41L2_HUMAN	Band 4.1-like protein 2 OS=Homo sapiens GN=EPB41L2 PE=1 SV=1	1	1	3.7	0.032		83.7	71.9	81.4	5032.8	6254.7	7062.4	3240.7	4630.7	2896.9	2168.0	1730.7	2261.4	452.6	263.7	532.3	205.7	154.3	126.3
SF3B2_HUMAN	Splicing factor 3B subunit 2 OS=Homo sapiens GN=SF3B2 PE=1 SV=2	1	1	3.7	0.125		99.7	141.0	119.5	26183.9	2226.3	34263.7	8833.0	10977.2	6552.0	3522.0	9663.7	9213.8	361.3	705.4	130.4	162.6	155.3	217.6
UGDH_HUMAN	UDP-glucose 6-dehydrogenase OS=Homo sapiens GN=UGDH PE=1 SV=1	1	1	3.7	0.021		2368.4	2366.3	2393.8	99135.9	93853.8	100047.0	129295.8	124400.6	134203.0	44061.2	51267.2	50746.0	13313.4	12295.5	11411.0	1081.7	1070.8	945.1
UBP5_HUMAN	Ubiquitin carboxyl-terminal hydrolase 5 OS=Homo sapiens GN=USP5 PE=1 SV=2	5	5	3.6	0.052		1335.7	1485.6	1449.6	131512.0	151094.2	138678.3	49616.8	55325.2	66031.3	48029.5	56196.1	47708.4	3169.7	4745.3	3350.6	516.2	484.6	446.6
MSH6_HUMAN	DNA mismatch repair protein Msh6 OS=Homo sapiens GN=MSH6 PE=1 SV=2	1	1	3.6	0.202		7.6	50.3	186.3	3018.4	6293.6	4891.2	204.8	826.6	70.7	0.0	4161.0	145.4	0.0	0.0	0.0	7.1	6.1	0.0
BASI_HUMAN	Basigin OS=Homo sapiens GN=BSG PE=1 SV=2	2	2	3.6	0.131		780.4	182.6	669.4	1439.2	2297.9	3683.6	19409.0	25126.0	27698.6	1915.3	5274.6	1927.2	3498.7	2974.7	4518.5	868.2	635.8	1138.4
HSPB1_HUMAN	Heat shock protein beta-1 OS=Homo sapiens GN=HSPB1 PE=1 SV=2	3	3	3.5	0.060		76.2	107.1	79.0	6668.4	14598.1	6716.0	6000.4	6213.6	3252.9	2054.3	2259.1	2857.7	1257.4	2303.9	1240.4	268.2	208.9	177.6
RNH2A_HUMAN	Ribonuclease H2 subunit A OS=Homo sapiens GN=RNASEH2A PE=1 SV=2	1	1	3.5	0.046		3317.2	3291.5	3211.7	286471.8	341682.6	356137.1	125426.1	157885.6	164269.9	90189.0	113597.0	101971.8	30061.0	37208.6	34308.0	3058.4	3074.2	3030.5
1433B_HUMAN	14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB PE=1 SV=3	10	7	3.5	0.065		445.5	461.6	406.3	31065.2	38218.3	8728.8	14320.8	17978.6	12138.8	8719.3	7096.9	6715.9	3494.0	4439.5	3361.0	747.6	676.7	558.0
DBNL_HUMAN	Drebrin-like protein OS=Homo sapiens GN=DBNL PE=1 SV=1	5	2	3.4	0.178		248.8	294.2	232.9	4244.0	2562.3	1893.5	23445.7	13565.9	3680.7	1586.4	5871.5	2634.2	765.6	1732.7	1025.7	309.3	451.0	366.0
PRS6B_HUMAN	26S protease regulatory subunit 6B OS=Homo sapiens GN=PSMC4 PE=1 SV=2	2	2	3.3	0.147		577.5	691.7	547.2	600.0	2453.1	481.1	15470.9	15103.0	18040.3	1766.4	600.8	658.2	4205.2	4313.1	3603.2	315.6	431.3	301.4
GLU2B_HUMAN	Glucosidase 2 subunit beta OS=Homo sapiens GN=PRKCSH PE=1 SV=2	9	6	3.3	0.056		4937.9	5223.4	4739.4	541836.5	558810.9	618485.8	241896.9	272853.2	206001.3	189142.3	189011.5	193368.4	59675.9	50626.4	48422.4	7723.6	7442.6	7293.4
GBLP_HUMAN	Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Homo sapiens GN=GNB2L1 PE=1 SV=3	6	6	3.3	0.070		918.2	701.3	716.3	42533.3	48456.9	30032.2	13186.1	13772.5	14903.8	11400.6	9716.5	9062.0	3819.0	4524.6	2185.3	3486.4	3424.6	3171.1
CAN1_HUMAN	Calpain-1 catalytic subunit OS=Homo sapiens GN=CAPN1 PE=1 SV=1	2	1	3.2	0.070		12.8	20.6	14.7	5901.9	6996.7	10005.6	18335.3	16660.6	21063.7	7117.7	6838.2	9154.9	625.8	325.8	114.8	59.2	67.9	63.8
DCTN1_HUMAN	Dynactin subunit 1 OS=Homo sapiens GN=DCTN1 PE=1 SV=3	4	1	3.2	0.067		617.4	769.5	708.5	157977.9	155615.9	170555.5	53910.5	69227.3	71875.5	67747.6	69928.8	57793.8	5201.0	4336.4	4660.7	1025.6	930.9	859.4
ANM1_HUMAN	Protein arginine N-methyltransferase 1 OS=Homo sapiens GN=PRMT1 PE=1 SV=2	3	2	3.2	0.089		489.8	356.2	379.5	27862.5	46666.5	9053.0	14046.1	11254.8	23105.2	5101.5	10705.8	11500.4	3386.7	5723.7	4056.6	451.5	398.1	415.6
PRS6A_HUMAN	26S protease regulatory subunit 6A OS=Homo sapiens GN=PSMC3 PE=1 SV=3	3	3	3.2	0.194		1301.0	1178.3	1037.2	0.0	0.0	164.6	24887.9	29762.6	30162.4	200.1	467.6	0.0	7744.4	9685.2	6915.9	782.0	968.6	1055.9
CPNE3_HUMAN	Copine-3 OS=Homo sapiens GN=CPNE3 PE=1 SV=1	12	10	3.1	0.051		24210.1	26001.4	25240.9	1796874.8	1807339.8	1878899.6	752341.1	860614.8	879715.3	522903.8	627088.7	589675.7	208968.0	276194.4	254287.2	31469.2	32620.8	31257.4
AXA2L_HUMAN	Putative annexin A2-like protein OS=Homo sapiens GN=ANXA2P2 PE=5 SV=2	21	1	3.1	0.052		364.5	370.8	353.1	25682.0	26945.5	29588.3	11971.8	14214.0	11120.3	7931.4	6874.7	7627.8	4798.4	5348.1	4732.9	555.6	517.2	532.2
SEPT2_HUMAN	Septin-2 OS=Homo sapiens GN=SEPT2 PE=1 SV=1	3	3	3.1	0.120		362.4	452.9	443.8	8130.6	5977.7	3512.5	7197.8	841.9	273.5	63.9	3844.3	3818.6	84.3	329.2	0.0	199.8	269.9	242.5
IPO5_HUMAN	Importin-5 OS=Homo sapiens GN=IPO5 PE=1 SV=4	5	4	3.0	0.079		1202.2	1482.3	1795.9	288829.9	143899.4	296938.5	143628.1	119518.0	107302.1	127957.8	145434.7	74033.1	3879.2	6317.5	5456.3	918.3	932.5	796.4
HSP7C_HUMAN	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1	91	37	3.0	0.050		87902.4	88787.8	93007.5	10289825.0	10363534.5	10155758.0	4998118.7	5546374.7	5302957.6	3736827.4	4088983.4	3892005.6	1045050.2	1229039.4	1082166.1	155833.8	155258.7	156515.3
FUBP3_HUMAN	Far upstream element-binding protein 3 OS=Homo sapiens GN=FUBP3 PE=1 SV=2	3	1	3.0	0.052		4493.4	4848.0	4113.7	759834.3	726769.7	642883.5	382178.3	415585.5	291440.4	218074.1	253931.6	217876.3	124058.7	118385.7	116852.8	9699.5	9521.9	10221.9
IF2B3_HUMAN	Insulin-like growth factor 2 mRNA-binding protein 3 OS=Homo sapiens GN=IGF2BP3 PE=1 SV=2	4	1	2.9	0.120		47.5	220.8	260.2	23173.5	18468.0	16926.0	8076.1	2818.1	3557.1	4970.0	6946.2	4213.9	2537.4	3686.9	2344.2	388.6	313.3	193.7
ALDOC_HUMAN	Fructose-bisphosphate aldolase C OS=Homo sapiens GN=ALDOC PE=1 SV=2	2	2	2.9	0.068		885.3	783.6	985.1	1649.6	0.0	3.3	1178.5	388.5	0.0	506.0	319.0	0.0	39.2	0.0	150.3	425.2	285.4	322.1
RL5_HUMAN	60S ribosomal protein L5 OS=Homo sapiens GN=RPL5 PE=1 SV=3	11	7	2.9	0.123		3360.8	3786.3	3513.0	180633.5	47437.9	147091.9	44901.7	44062.2	39482.6	19616.4	16441.1	21414.3	34366.6	37377.0	30350.0	6741.5	7534.5	6435.9
GTF2I_HUMAN	General transcription factor II-I OS=Homo sapiens GN=GTF2I PE=1 SV=2	1	1	2.8	0.048		27390.5	30322.4	27862.4	2135943.2	2211887.7	2233473.4	1130841.3	1369823.6	1317563.7	762966.8	828533.5	807093.6	355898.1	475039.4	390083.4	32790.4	37149.5	32242.4
G0S2_HUMAN	G0/G1 switch protein 2 OS=Homo sapiens GN=G0S2 PE=1 SV=1	2	2	2.8	0.074		8539.4	8616.3	9049.8	7814.7	3015.3	1795.5	28755.1	30226.7	4885.3	1774.2	3012.5	4192.9	3633.5	1809.3	781.1	7022.0	7315.1	7063.6
RUVB2_HUMAN	RuvB-like 2 OS=Homo sapiens GN=RUVBL2 PE=1 SV=3	12	10	2.8	0.079		6177.5	6049.2	5949.8	36137.8	46256.3	44226.4	123321.3	131746.0	143212.9	18138.1	33098.8	23135.4	37389.3	31911.0	34957.4	5545.9	5927.8	4942.0
GRP75_HUMAN	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2	40	35	2.8	0.062		34923.2	36015.2	35905.9	2446864.9	2506791.7	2375293.8	1265453.5	1470539.6	1201023.5	1110175.1	1237344.5	1112090.9	168184.3	187649.9	168905.1	44361.1	44270.0	42861.2
FEN1_HUMAN	Flap endonuclease 1 OS=Homo sapiens GN=FEN1 PE=1 SV=1	3	3	2.7	0.081		1285.9	1221.1	1151.9	38823.0	765.7	34519.9	46392.0	52378.8	52331.0	13149.4	16133.9	11846.7	13018.8	13346.2	12894.8	1412.1	1449.7	1323.6
PABP4_HUMAN	Polyadenylate-binding protein 4 OS=Homo sapiens GN=PABPC4 PE=1 SV=1	13	3	2.7	0.060		1854.5	1983.9	1911.5	70072.4	87634.5	87734.1	48571.5	67195.1	56567.8	46672.4	53198.6	46756.8	2091.7	1605.2	1235.0	2224.5	2288.9	2271.4
NDUS1_HUMAN	NADH-ubiquinone oxidoreductase 75 kDa subunit, mitochondrial OS=Homo sapiens GN=NDUFS1 PE=1 SV=3	2	1	2.7	0.169		103.6	100.1	35.5	6705.6	9543.2	3432.2	0.0	4048.8	1672.8	2403.1	1780.6	2927.8	714.5	389.7	1173.1	59.5	96.1	82.1
NYNRI_HUMAN	Protein NYNRIN OS=Homo sapiens GN=NYNRIN PE=2 SV=3	1	1	2.7	0.066		3162.3	3422.8	2946.0	343152.8	287439.9	229817.7	231972.4	200458.8	186891.6	160873.7	196156.1	164214.0	7890.4	10210.7	6373.1	5103.7	4485.2	4749.8
GFAP_HUMAN	Glial fibrillary acidic protein OS=Homo sapiens GN=GFAP PE=1 SV=1	4	2	2.7	0.126		1296.1	1112.4	1166.3	7939.8	5707.2	6100.7	29728.7	30674.4	29461.9	2726.8	2558.0	2147.5	11108.7	12518.7	8950.6	602.8	1026.2	985.4
RL4_HUMAN	60S ribosomal protein L4 OS=Homo sapiens GN=RPL4 PE=1 SV=5	31	28	2.6	0.154		31823.3	32875.8	32908.0	117824.3	114517.2	165041.8	804478.9	926904.7	912243.9	79801.3	97819.3	99360.5	234317.9	300594.6	277493.7	32985.6	35573.3	33884.8
CORO7_HUMAN	Coronin-7 OS=Homo sapiens GN=CORO7 PE=1 SV=2	3	2	2.6	0.070		700.9	1014.0	580.2	38208.4	63308.9	53580.4	24797.4	27198.8	28596.9	11265.9	19471.8	9585.5	13540.0	17167.0	15969.9	1099.2	1193.9	1180.5
DNM1L_HUMAN	Dynamin-1-like protein OS=Homo sapiens GN=DNM1L PE=1 SV=2	4	3	2.6	0.084		400.6	419.1	403.3	26962.2	39228.6	25279.5	18861.3	31878.2	13047.7	22696.9	16753.7	16363.4	1069.9	1004.1	225.5	517.6	439.4	472.9
HNRH1_HUMAN	Heterogeneous nuclear ribonucleoprotein H OS=Homo sapiens GN=HNRNPH1 PE=1 SV=4	9	5	2.6	0.063		10452.3	9733.4	11794.3	328966.9	349225.4	340036.1	153900.2	162841.5	167313.8	108269.1	132758.5	112164.5	64888.7	76612.3	66321.6	8295.5	8266.9	8806.5
MCM3_HUMAN	DNA replication licensing factor MCM3 OS=Homo sapiens GN=MCM3 PE=1 SV=3	12	11	2.6	0.057		5956.5	6295.3	6334.8	455468.5	447102.8	358823.1	286321.9	250650.7	281643.2	200525.9	222261.0	175175.2	56246.4	72516.3	57143.2	6657.3	5937.4	6685.7
OST48_HUMAN	olichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit OS=Homo sapiens GN=DDOST PE=1 SV	5	4	2.6	0.231		2375.3	2594.1	3002.8	1538.4	1098.8	360.2	74204.3	89358.4	69342.0	6848.2	9479.9	6504.0	19055.8	20481.7	18749.6	4298.9	3843.6	4247.4
ALBU_HUMAN	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	33	27	2.6	0.052		122683.5	123691.6	129828.2	6786366.8	6922916.9	6881983.0	4032439.2	4602727.1	4805631.1	3079946.7	3338904.3	3105926.4	894767.1	1258012.7	1016003.2	170314.3	174426.2	175102.3
HMGB2_HUMAN	High mobility group protein B2 OS=Homo sapiens GN=HMGB2 PE=1 SV=2	5	2	2.6	0.212		38.1	56.9	8.1	1037.1	5975.5	249.2	2854.2	1655.3	1705.1	360.7	2270.8	120.6	519.8	183.4	1508.8	78.9	133.5	58.2
SEPT7_HUMAN	Septin-7 OS=Homo sapiens GN=SEPT7 PE=1 SV=2	7	6	2.6	0.051		4672.5	4817.4	5096.2	74005.3	118886.7	127893.2	158637.2	172497.3	169714.2	67558.9	63844.7	64145.2	39853.1	34872.1	39976.5	3931.2	4417.8	3940.2
METK2_HUMAN	S-adenosylmethionine synthase isoform type-2 OS=Homo sapiens GN=MAT2A PE=1 SV=1	4	3	2.6	0.141		2548.8	2699.3	2877.7	9433.0	4789.8	22528.3	54428.6	72490.0	61069.5	10339.4	8359.8	7049.0	18347.1	20321.4	17535.0	2347.6	2878.5	2770.7
KAD2_HUMAN	Adenylate kinase 2, mitochondrial OS=Homo sapiens GN=AK2 PE=1 SV=2	4	4	2.6	0.192		423.4	489.3	485.3	88326.6	75092.9	5848.0	6344.0	38743.4	55317.2	3572.6	49399.4	41474.7	1851.9	2280.8	1305.4	1702.9	1444.9	1734.4
MCM4_HUMAN	DNA replication licensing factor MCM4 OS=Homo sapiens GN=MCM4 PE=1 SV=5	11	8	2.6	0.072		4317.5	4360.1	4858.6	432591.7	433993.1	182115.4	244308.4	268868.3	261962.6	182555.6	178515.4	188879.0	45406.3	60841.6	44379.6	3977.4	3888.1	4017.8
STML2_HUMAN	Stomatin-like protein 2, mitochondrial OS=Homo sapiens GN=STOML2 PE=1 SV=1	3	2	2.5	0.189		734.6	622.1	922.1	3094.4	0.0	0.0	331.5	2211.2	0.0	453.9	0.0	0.0	889.9	0.0	86.0	606.4	502.3	566.2
GRP78_HUMAN	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2	53	41	2.5	0.065		105580.9	107857.7	112831.0	7778907.3	7191777.9	7162221.5	3903704.9	4185372.8	3974908.7	2874842.9	3251635.9	2792322.5	1159015.9	1498889.8	1365983.7	202312.9	197995.9	203494.3
PSMD1_HUMAN	26S proteasome non-ATPase regulatory subunit 1 OS=Homo sapiens GN=PSMD1 PE=1 SV=2	7	5	2.5	0.058		44844.1	46700.8	56687.4	4977758.6	5548752.0	6027978.2	3212913.0	3216802.7	4359517.5	2038576.0	2141054.0	2231085.1	1061422.4	1233636.1	1679408.3	139663.0	137221.1	152828.2
SYMC_HUMAN	Methionine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=MARS PE=1 SV=2	9	6	2.5	0.062		3308.3	3085.2	3412.8	289219.7	186494.7	207538.1	153522.3	153362.2	184879.7	120821.8	128918.5	82695.2	63743.5	8739.1	54343.8	2644.8	2259.5	2553.3
STIP1_HUMAN	Stress-induced-phosphoprotein 1 OS=Homo sapiens GN=STIP1 PE=1 SV=1	35	30	2.5	0.056		25624.1	26083.0	25021.0	1378939.6	1444925.3	1411212.2	927207.9	1026433.7	985406.3	698700.2	760440.0	695682.8	185946.1	234859.8	204709.2	28375.0	28961.5	28185.3
SF3B3_HUMAN	Splicing factor 3B subunit 3 OS=Homo sapiens GN=SF3B3 PE=1 SV=4	7	6	2.5	0.087		2628.2	2396.2	2196.2	33595.8	81156.8	118986.7	36084.2	63561.6	74248.9	36167.3	44407.8	29815.6	12887.7	17574.5	17777.7	1942.5	1829.8	1732.1
SNX2_HUMAN	Sorting nexin-2 OS=Homo sapiens GN=SNX2 PE=1 SV=2	4	3	2.5	0.142		621.4	773.4	871.0	127876.5	122190.3	13467.6	36826.6	84477.1	104044.3	34260.4	76393.6	72048.8	3039.6	3100.2	3242.9	927.7	678.3	853.5
MOES_HUMAN	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3	23	9	2.5	0.077		4218.7	4452.0	4351.7	311940.0	399329.6	359743.7	168872.7	197266.9	181009.6	146074.0	171913.6	151599.1	46810.1	62284.9	47200.2	6809.0	7144.2	6378.1
ATPB_HUMAN	ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3	27	26	2.5	0.206		52891.4	52745.5	54603.2	82788.6	82377.4	103751.1	1325200.2	1459946.5	1414346.4	144332.7	169453.9	162248.1	376197.5	424923.0	386978.2	58164.6	54507.3	58275.4
LRP1_HHV1F	Latency-related protein 1 OS=Human herpesvirus 1 (strain F) PE=4 SV=1	1	1	2.5	0.284		2681.2	2635.1	2401.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	225.3	0.0	0.0	0.0	0.0	1005.0	864.5	974.2
COPB_HUMAN	Coatomer subunit beta OS=Homo sapiens GN=COPB1 PE=1 SV=3	10	7	2.5	0.089		1346.4	1568.3	1673.1	188362.8	167932.9	208570.0	115059.0	124165.1	133637.6	136655.8	120016.6	111413.7	3035.2	3228.2	2182.2	1049.7	1034.8	1227.5
QCR1_HUMAN	Cytochrome b-c1 complex subunit 1, mitochondrial OS=Homo sapiens GN=UQCRC1 PE=1 SV=3	1	1	2.5	0.060		696.0	615.9	593.3	584.2	540.5	0.0	238.7	1220.8	1951.9	215.9	210.1	0.0	315.5	722.4	186.0	240.3	402.6	309.7
RBM39_HUMAN	RNA-binding protein 39 OS=Homo sapiens GN=RBM39 PE=1 SV=2	2	1	2.5	0.079		266.7	294.3	288.9	19278.8	23820.4	19368.8	20519.4	18008.8	28547.8	15005.1	18014.3	16540.3	1007.1	965.9	970.1	96.5	90.0	123.2
IQGA1_HUMAN	Ras GTPase-activating-like protein IQGAP1 OS=Homo sapiens GN=IQGAP1 PE=1 SV=1	12	10	2.5	0.062		5032.3	5128.7	4885.7	194414.9	197882.9	204875.6	221755.5	240357.9	238267.5	150345.2	165423.2	142753.2	24198.7	22757.2	19109.0	2277.6	2473.0	2465.8
BIEA_HUMAN	Biliverdin reductase A OS=Homo sapiens GN=BLVRA PE=1 SV=2	3	3	2.5	0.139		848.2	791.5	813.8	2036.3	3736.2	3589.7	9297.0	6213.2	16337.3	3427.6	2330.5	2132.8	3993.5	2198.3	1846.5	595.3	628.9	616.2
1433T_HUMAN	14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1 SV=1	8	3	2.4	0.444		3.6	13.9	14.8	1130.1	453.4	0.0	0.0	0.0	0.0	295.8	95.2	0.0	135.0	100.5	0.0	23.6	4.7	4.7
MPCP_HUMAN	Phosphate carrier protein, mitochondrial OS=Homo sapiens GN=SLC25A3 PE=1 SV=2	8	5	2.4	0.047		5250.1	5102.8	4721.8	165462.9	182135.2	194550.2	142400.4	153248.1	152516.8	81986.9	95266.0	83353.1	40634.4	44927.8	41552.8	7233.1	7570.0	8355.3
PLST_HUMAN	Plastin-3 OS=Homo sapiens GN=PLS3 PE=1 SV=4	43	27	2.4	0.063		38337.3	38211.1	39111.3	1872929.1	2114075.6	2271486.3	1506358.7	1661760.9	1575040.9	1150430.5	1255153.7	1216682.0	239712.0	290862.9	277164.4	45632.0	43038.0	44527.6
KPYM_HUMAN	Pyruvate kinase PKM OS=Homo sapiens GN=PKM PE=1 SV=4	57	48	2.4	0.072		128457.5	129400.9	129222.5	9373523.0	9461380.8	8512027.9	5032900.0	6045160.8	5955413.6	4338981.6	5077801.5	4703304.7	1194072.7	1383186.2	1323899.6	140591.9	142167.2	140997.8
PUF60_HUMAN	Poly(U)-binding-splicing factor PUF60 OS=Homo sapiens GN=PUF60 PE=1 SV=1	3	3	2.4	0.063		1383.9	1777.8	1884.9	83924.0	68085.7	89414.5	96439.7	86556.0	79634.3	49671.8	62969.4	58373.1	14211.2	14079.4	7686.2	1085.3	969.7	932.9
ANXA5_HUMAN	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2	17	14	2.4	0.057		3656.8	3452.9	3704.9	63099.3	91475.9	90999.2	44910.7	49952.6	58691.8	33150.4	34317.0	33269.0	11292.1	14070.3	9018.5	11456.5	11432.2	11557.5
LA_HUMAN	Lupus La protein OS=Homo sapiens GN=SSB PE=1 SV=2	3	3	2.4	0.181		5009.0	4687.0	5070.0	18973.2	28999.1	34773.9	174359.5	201552.0	170799.5	31493.3	36429.9	23288.9	50330.5	55538.6	54848.1	5196.8	4972.3	5345.8
SET_HUMAN	Protein SET OS=Homo sapiens GN=SET PE=1 SV=3	12	2	2.4	0.034		2041.8	2415.1	2131.4	18204.2	17036.8	27545.5	16942.2	13828.1	19739.5	5569.3	7108.3	4807.5	7143.4	8050.4	5534.1	3800.6	4096.6	3774.0
ATP4A_HUMAN	Potassium-transporting ATPase alpha chain 1 OS=Homo sapiens GN=ATP4A PE=2 SV=5	2	1	2.4	0.065		154786.0	153693.4	150087.0	16548798.2	16382320.5	17077817.4	10163613.0	11133042.9	11096905.1	6466090.4	7405739.2	6784326.3	3664389.8	4570723.9	4450784.7	409632.5	401236.6	401283.2
RL6_HUMAN	60S ribosomal protein L6 OS=Homo sapiens GN=RPL6 PE=1 SV=3	18	17	2.4	0.043		15140.9	14722.2	13023.9	57079.9	107563.5	86125.2	60899.3	89409.0	150060.5	44317.6	48933.8	38264.1	23229.5	25507.7	27381.2	13312.0	14103.4	13299.9
XPO2_HUMAN	Exportin-2 OS=Homo sapiens GN=CSE1L PE=1 SV=3	28	21	2.4	0.049		90968.9	90027.8	89118.2	1246784.1	1156314.5	1120728.6	735861.4	773816.2	832812.4	635206.6	648073.8	623167.9	99550.0	133758.5	116343.2	98832.6	104591.8	108432.1
TRFL_HUMAN	Lactotransferrin OS=Homo sapiens GN=LTF PE=1 SV=6	1	1	2.4	0.099		223.8	184.8	178.8	9049.7	19719.1	22953.2	11399.9	17632.7	15178.5	10489.6	12976.6	12237.7	820.3	2046.8	1076.9	323.8	260.5	275.0
RFA1_HUMAN	Replication protein A 70 kDa DNA-binding subunit OS=Homo sapiens GN=RPA1 PE=1 SV=2	4	4	2.4	0.090		1233.6	1542.7	1454.2	76808.6	78666.0	72605.6	111251.9	138807.7	141172.7	68025.6	79039.5	71322.9	8575.4	15868.1	14137.9	1658.1	1729.3	1435.1
PYRG1_HUMAN	CTP synthase 1 OS=Homo sapiens GN=CTPS1 PE=1 SV=2	2	2	2.4	0.060		1325.9	1253.8	1135.6	49033.0	53271.7	42479.3	44257.0	55246.8	49336.4	31572.9	29078.6	32638.7	10490.2	7693.8	8385.9	2103.3	1908.5	2087.7
CNDP2_HUMAN	Cytosolic non-specific dipeptidase OS=Homo sapiens GN=CNDP2 PE=1 SV=2	2	2	2.4	0.251		437.2	383.2	465.9	1302.3	845.7	1494.6	14582.3	19456.0	21462.8	7562.1	7032.0	5601.2	1414.7	1762.5	935.1	431.9	442.8	462.0
ENOA_HUMAN	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	70	58	2.3	0.197		261248.8	264549.0	264882.6	1241570.6	1460263.5	1600666.2	8877020.2	10277094.9	10086478.9	817614.1	1157766.0	1014897.6	3034393.9	3963883.3	3672210.2	334277.7	335770.4	344787.6
K22O_HUMAN	Keratin, type II cytoskeletal 2 oral OS=Homo sapiens GN=KRT76 PE=1 SV=2	17	9	2.3	0.071		16861.3	18526.6	17762.6	1390580.3	1506236.0	1532124.8	819281.4	963503.9	964669.6	495447.0	577129.8	551068.3	404155.6	500540.2	444771.9	39618.5	46160.1	42823.0
TKT_HUMAN	Transketolase OS=Homo sapiens GN=TKT PE=1 SV=3	28	23	2.3	0.069		15362.3	16222.6	16249.0	1022828.4	1019868.5	1018222.8	915205.0	999356.5	934623.7	667267.0	676573.6	683540.7	150496.0	181497.1	164594.5	15385.0	14460.6	13608.2
UBP14_HUMAN	Ubiquitin carboxyl-terminal hydrolase 14 OS=Homo sapiens GN=USP14 PE=1 SV=3	3	3	2.3	0.112		5947.6	4920.6	5137.9	489236.4	563664.9	254107.9	186175.9	277755.5	215735.7	211909.9	145767.6	181479.6	111793.1	113194.8	71388.5	9432.6	10074.7	8402.2
PUR2_HUMAN	Trifunctional purine biosynthetic protein adenosine-3 OS=Homo sapiens GN=GART PE=1 SV=1	9	6	2.3	0.050		3844.6	3818.9	3763.3	95316.8	87887.0	70237.4	93309.2	88581.4	90427.7	46524.1	53212.7	35117.9	19007.9	34754.0	24994.4	6041.3	5927.2	6223.9
PABP1_HUMAN	Polyadenylate-binding protein 1 OS=Homo sapiens GN=PABPC1 PE=1 SV=2	28	7	2.3	0.069		4602.9	5340.2	4820.3	276215.7	283837.8	285838.7	172559.0	195833.0	201780.9	117023.1	152982.6	135420.7	52824.9	68413.2	66537.4	8503.7	8608.2	8919.3
PHB_HUMAN	Prohibitin OS=Homo sapiens GN=PHB PE=1 SV=1	8	6	2.3	0.061		1327.1	1582.4	1370.0	53605.1	67251.7	78619.9	51239.4	60736.3	57913.0	28896.5	35096.2	31403.1	17647.3	22697.8	20331.7	1868.9	2069.8	1846.7
MYH10_HUMAN	Myosin-10 OS=Homo sapiens GN=MYH10 PE=1 SV=3	31	17	2.3	0.072		7170.6	6918.0	6885.5	451394.0	509579.7	482524.0	263612.5	336405.0	314654.8	194957.9	174752.1	173327.0	146625.5	164542.8	149916.5	10871.0	10940.7	10407.8
NP1L4_HUMAN	Nucleosome assembly protein 1-like 4 OS=Homo sapiens GN=NAP1L4 PE=1 SV=1	7	3	2.3	0.078		1516.2	1874.7	2158.9	61630.4	71627.0	88214.5	71595.4	86198.4	90517.0	59701.6	57620.9	52163.6	11658.3	12554.8	10018.8	1436.7	1553.2	1281.2
XRCC6_HUMAN	X-ray repair cross-complementing protein 6 OS=Homo sapiens GN=XRCC6 PE=1 SV=2	22	18	2.3	0.084		20371.9	20311.1	19734.9	1271709.9	1391206.5	1318983.5	1104444.8	1245865.4	1205780.4	863466.8	1011696.9	985580.6	109440.7	166122.8	148522.7	22548.8	23349.7	22185.5
SMC2_HUMAN	Structural maintenance of chromosomes protein 2 OS=Homo sapiens GN=SMC2 PE=1 SV=2	5	2	2.3	0.079		3926.8	3923.4	3825.7	188940.0	208469.4	196007.0	148805.9	148363.9	146733.5	110536.8	131617.2	123632.7	28620.5	30503.6	28625.4	3290.1	3265.3	3138.0
SRSF6_HUMAN	Serine/arginine-rich splicing factor 6 OS=Homo sapiens GN=SRSF6 PE=1 SV=2	3	3	2.3	0.086		645.5	322.7	196.2	12824.2	5946.4	4756.4	13213.3	11052.2	9863.7	3083.3	3901.3	2403.3	5189.9	4134.2	5486.0	621.5	774.3	426.2
TPM3_HUMAN	Tropomyosin alpha-3 chain OS=Homo sapiens GN=TPM3 PE=1 SV=2	2	2	2.3	0.277		298.8	159.7	235.7	319.1	823.5	3378.3	6951.2	3835.0	13068.1	2119.3	4020.0	4698.5	651.4	87.9	410.9	344.2	268.8	290.0
HNRPF_HUMAN	Heterogeneous nuclear ribonucleoprotein F OS=Homo sapiens GN=HNRNPF PE=1 SV=3	10	9	2.3	0.209		9977.6	11098.5	9412.1	30691.3	43028.5	46269.6	247508.3	288676.1	289384.0	24684.2	21529.9	18994.1	103237.4	115324.4	114988.5	11027.6	11726.8	11338.5
YBOX3_HUMAN	Y-box-binding protein 3 OS=Homo sapiens GN=YBX3 PE=1 SV=4	6	3	2.3	0.087		1806.3	2569.2	2621.4	70073.7	86314.5	103021.1	105084.6	108099.9	123087.0	68378.1	78970.5	73122.0	13093.6	13307.4	14762.9	2115.4	1917.8	1920.8
MATR3_HUMAN	Matrin-3 OS=Homo sapiens GN=MATR3 PE=1 SV=2	14	10	2.2	0.069		5364.7	5197.1	5454.3	802833.6	744136.1	771374.4	568841.3	570508.4	602151.0	307182.9	316895.8	312887.1	244134.4	305697.5	261880.6	21620.3	22051.3	22575.3
BLMH_HUMAN	Bleomycin hydrolase OS=Homo sapiens GN=BLMH PE=1 SV=1	2	2	2.2	0.059		365.4	334.8	285.3	9482.5	11266.3	9592.0	10800.7	10924.5	9884.4	3959.0	5042.6	3286.5	4458.1	6193.1	4207.9	279.0	377.7	234.8
PDIA4_HUMAN	Protein disulfide-isomerase A4 OS=Homo sapiens GN=PDIA4 PE=1 SV=2	16	13	2.2	0.077		7935.7	8352.1	8221.6	700490.2	708490.7	697850.8	405804.4	466086.5	483098.3	271343.6	327486.8	298148.0	173441.7	228573.1	186856.3	24745.5	21722.8	22716.7
YBOX1_HUMAN	Nuclease-sensitive element-binding protein 1 OS=Homo sapiens GN=YBX1 PE=1 SV=3	5	3	2.2	0.280		7111.6	7810.4	7125.5	0.0	692.8	0.0	101948.0	108364.3	107110.2	1433.4	1885.5	2790.9	44030.0	46369.4	40071.0	4987.1	5514.7	5298.3
RS3_HUMAN	40S ribosomal protein S3 OS=Homo sapiens GN=RPS3 PE=1 SV=2	11	7	2.2	0.078		745.1	862.1	712.0	12636.6	28822.7	14122.6	16168.2	21180.6	20861.1	10793.3	7915.1	12144.8	4585.7	4223.1	4468.0	3046.3	2398.3	2590.8
DLDH_HUMAN	Dihydrolipoyl dehydrogenase, mitochondrial OS=Homo sapiens GN=DLD PE=1 SV=2	2	1	2.2	0.191		0.0	0.0	0.0	10960.2	11562.2	11999.7	1210.4	3909.4	6524.1	6357.8	6815.5	6452.4	386.6	275.0	469.7	19.3	13.4	20.7
RPN1_HUMAN	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPN1 PE=1 SV=1	15	13	2.2	0.077		125230.3	127682.5	126706.1	14203160.3	13613942.6	13836372.7	9015439.4	10253483.8	10243856.6	6053961.5	6896998.0	6568986.0	3012799.8	4802279.4	3911193.4	330516.2	352661.6	343388.4
TXND5_HUMAN	Thioredoxin domain-containing protein 5 OS=Homo sapiens GN=TXNDC5 PE=1 SV=2	5	3	2.2	0.082		102934.4	101062.7	112899.4	11775127.0	11383598.4	11620406.6	7223409.8	7082153.4	8266481.9	4846164.3	5024246.8	4108708.1	3252640.4	4092722.5	3904589.0	349522.6	328687.9	344591.2
IMA1_HUMAN	Importin subunit alpha-1 OS=Homo sapiens GN=KPNA2 PE=1 SV=1	7	5	2.2	0.103		2177.2	2146.6	2133.2	106389.9	125559.8	86413.9	75985.6	99368.8	110102.0	70851.9	83864.7	86934.2	8848.4	14831.6	9428.1	1104.1	999.8	1157.4
PAIRB_HUMAN	Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERBP1 PE=1 SV=2	9	7	2.2	0.068		12058.2	12131.6	12522.4	454943.0	434929.3	455070.5	446163.9	502499.2	484205.5	250254.3	297544.5	283391.4	125671.0	149053.3	141037.1	12438.4	13501.8	10774.2
DDX3X_HUMAN	ATP-dependent RNA helicase DDX3X OS=Homo sapiens GN=DDX3X PE=1 SV=3	13	8	2.2	0.071		4870.1	4143.9	5222.8	574709.3	479704.5	548377.7	457846.6	534598.3	582562.5	218792.7	159191.5	251764.5	248159.9	258621.0	254631.9	26443.6	22729.3	24519.3
TCPQ_HUMAN	T-complex protein 1 subunit theta OS=Homo sapiens GN=CCT8 PE=1 SV=4	39	29	2.2	0.081		33926.7	34187.9	33846.3	1602511.4	1590507.4	1538156.6	1141330.4	1485234.8	1326649.0	904166.2	1086870.1	958210.8	300291.0	384575.4	309713.5	34018.9	31644.9	32659.6
SYRC_HUMAN	Arginine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=RARS PE=1 SV=2	1	1	2.2	0.098		1021.2	912.1	319.6	124593.7	105249.3	60694.8	94283.9	130242.6	76936.0	57789.7	71501.5	55388.3	30411.5	28272.3	23395.7	2545.3	2359.0	2110.3
RS8_HUMAN	40S ribosomal protein S8 OS=Homo sapiens GN=RPS8 PE=1 SV=2	5	3	2.2	0.357		118.3	106.9	75.7	2388.1	5279.2	30664.2	4729.9	4139.2	5576.6	4273.2	4499.0	3234.2	4557.8	4948.5	1895.6	388.8	324.1	474.4
KATL2_HUMAN	Katanin p60 ATPase-containing subunit A-like 2 OS=Homo sapiens GN=KATNAL2 PE=2 SV=3	4	2	2.2	0.078		10164.8	10545.3	11217.2	1097398.4	1048841.4	1085174.6	720514.1	788602.4	919930.4	406246.6	445943.4	437385.1	388344.2	401317.2	448030.3	39075.3	36903.7	38904.5
IF2G_HUMAN	Eukaryotic translation initiation factor 2 subunit 3 OS=Homo sapiens GN=EIF2S3 PE=1 SV=3	9	4	2.2	0.136		15808.8	16110.0	15559.5	47225.6	10071.2	10525.1	101898.1	103354.7	127008.7	38509.7	29890.6	7780.4	29811.7	7584.2	28469.2	23102.6	20953.1	21780.9
VAT1_HUMAN	Synaptic vesicle membrane protein VAT-1 homolog OS=Homo sapiens GN=VAT1 PE=1 SV=2	3	2	2.1	0.097		15395.5	15663.1	15789.0	1558157.8	1569085.4	1608242.5	1453188.9	1715169.9	1567709.4	1178520.0	1329228.3	1088403.9	232804.4	270280.5	261736.6	27104.7	25698.7	26212.6
FUBP2_HUMAN	Far upstream element-binding protein 2 OS=Homo sapiens GN=KHSRP PE=1 SV=4	32	27	2.1	0.100		31473.4	32106.9	31613.3	2722835.3	2790534.1	2868705.9	1455445.3	1574013.8	1649839.3	1097189.9	1266632.5	1115765.6	686494.5	953027.4	779463.5	91915.0	91612.4	92026.5
KTN1_HUMAN	Kinectin OS=Homo sapiens GN=KTN1 PE=1 SV=1	6	3	2.1	0.086		2844.5	2846.2	2865.0	282874.8	234816.0	309568.9	195822.8	183988.1	211183.0	90264.4	125785.8	91239.3	100530.1	133394.4	101252.6	8324.3	9916.1	10522.2
CH60_HUMAN	60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2	51	47	2.1	0.095		133181.8	143451.4	102638.8	7244650.9	7465485.8	6919038.1	4400742.5	5246997.2	5023154.7	3870367.3	4363570.7	3985109.4	1263701.7	1750452.5	1490388.2	177651.8	173639.2	191960.6
PRDX4_HUMAN	Peroxiredoxin-4 OS=Homo sapiens GN=PRDX4 PE=1 SV=1	4	2	2.1	0.296		44.8	66.4	34.2	676.9	3314.8	0.0	5055.7	841.8	2783.2	2623.0	455.0	1847.3	641.3	77.4	233.3	60.2	67.3	31.1
TCPH_HUMAN	T-complex protein 1 subunit eta OS=Homo sapiens GN=CCT7 PE=1 SV=2	22	19	2.1	0.095		12680.8	13034.1	13439.1	598052.4	662197.2	622577.7	593163.5	383790.7	536964.9	407340.5	341294.0	449271.2	136818.4	170200.9	79164.9	12109.0	12492.5	12607.8
SRC8_HUMAN	Src substrate cortactin OS=Homo sapiens GN=CTTN PE=1 SV=2	10	9	2.1	0.074		25430.1	27202.3	29058.6	404327.3	473446.9	453800.5	233965.0	272974.8	288348.5	159285.3	176623.2	168022.0	133384.2	181142.5	146162.7	27455.9	26133.2	26242.2
DDX5_HUMAN	Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 PE=1 SV=1	16	8	2.1	0.131		1389.2	1335.6	1397.4	99526.9	152610.6	141661.2	114859.6	81119.1	106484.7	83748.0	101817.8	110132.4	11101.4	9677.7	8717.6	2373.6	1975.4	2000.1
2AAA_HUMAN	e/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha isoform OS=Homo sapiens GN=PPP2R1A PE=1	20	17	2.1	0.083		21395.4	21645.5	22829.8	506016.4	601102.4	494857.0	395627.0	439038.9	420253.7	307251.9	356980.7	326337.3	102669.5	120768.3	99425.4	25111.0	24504.5	25872.3
TCPE_HUMAN	T-complex protein 1 subunit epsilon OS=Homo sapiens GN=CCT5 PE=1 SV=1	23	18	2.1	0.098		15146.6	15354.2	15131.2	1017027.7	1016798.5	958382.2	571838.8	670377.0	600058.7	471279.5	510555.9	428021.3	244544.4	293473.0	295551.3	23808.8	27312.6	27542.6
TCPG_HUMAN	T-complex protein 1 subunit gamma OS=Homo sapiens GN=CCT3 PE=1 SV=4	21	18	2.1	0.097		12591.3	12897.4	11502.1	877301.7	959375.8	862952.9	739245.8	816617.5	802964.3	629490.9	620242.2	601102.3	149985.4	199948.5	171450.7	17526.8	18814.6	17384.5
LMNA_HUMAN	Prelamin-A/C OS=Homo sapiens GN=LMNA PE=1 SV=1	3	2	2.1	0.089		1062.2	1045.5	945.1	20798.6	35060.5	46355.1	33862.6	30449.1	35913.2	19312.7	19737.0	15119.7	14369.2	11764.0	13335.0	1379.7	1812.0	1195.6
TRAP1_HUMAN	Heat shock protein 75 kDa, mitochondrial OS=Homo sapiens GN=TRAP1 PE=1 SV=3	19	13	2.1	0.109		17249.2	16970.9	17713.5	493569.3	338967.8	224784.0	347567.7	268085.9	321354.9	255972.0	274871.5	247344.9	40987.6	38709.2	62139.5	19138.0	18669.7	19863.2
RS4X_HUMAN	40S ribosomal protein S4, X isoform OS=Homo sapiens GN=RPS4X PE=1 SV=2	7	1	2.1	0.423		62.6	53.7	50.6	0.0	0.0	0.0	136.1	43.1	484.0	0.0	0.0	0.0	253.2	0.0	82.8	46.5	6.6	8.0
ADT3_HUMAN	ADP/ATP translocase 3 OS=Homo sapiens GN=SLC25A6 PE=1 SV=4	9	1	2.1	0.088		211.9	191.2	224.1	14343.9	19061.9	17196.0	18194.2	15792.4	16187.2	11063.1	10634.0	10255.8	4946.5	5034.9	5468.1	348.2	431.4	349.0
FKBP4_HUMAN	Peptidyl-prolyl cis-trans isomerase FKBP4 OS=Homo sapiens GN=FKBP4 PE=1 SV=3	29	23	2.1	0.077		50407.9	50991.4	51929.9	831009.4	887037.9	842213.2	634766.3	764862.8	751414.0	522192.9	625180.0	547045.0	135280.2	184519.1	134489.7	63161.7	59332.1	60406.2
NONO_HUMAN	Non-POU domain-containing octamer-binding protein OS=Homo sapiens GN=NONO PE=1 SV=4	14	10	2.1	0.082		5072.0	5244.0	5348.4	252181.9	244643.4	260399.1	206194.4	235829.4	245989.2	133284.2	157003.4	152086.9	79990.5	88067.6	68205.0	7130.5	8291.4	7470.1
PRDX2_HUMAN	Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5	3	3	2.1	0.079		445.3	707.4	489.3	976.8	4544.1	2695.2	3213.5	4941.2	4728.1	1148.7	2061.2	1910.0	1439.1	1371.7	1190.5	614.3	530.4	669.9
IF2B1_HUMAN	Insulin-like growth factor 2 mRNA-binding protein 1 OS=Homo sapiens GN=IGF2BP1 PE=1 SV=2	19	14	2.1	0.095		6509.5	7427.5	7199.6	272598.8	293909.6	316473.0	332584.7	343048.1	250124.1	202981.3	222330.4	223199.0	69243.2	59156.0	76138.8	9188.5	9289.3	9434.1
CALX_HUMAN	Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2	15	13	2.1	0.162		5361.3	5434.5	5391.4	302109.8	227155.4	242934.6	537176.2	503982.4	481261.2	329337.3	356741.8	316141.1	21371.8	49915.8	20742.4	7358.7	7334.3	7387.0
RL13_HUMAN	60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4	8	8	2.1	0.105		669.2	634.6	628.1	23729.6	31453.6	39108.3	42094.2	31278.3	28610.9	31710.7	12615.9	15251.3	9424.7	13471.2	10823.4	979.3	979.7	873.8

EF1D_HUMAN	Elongation factor 1-delta OS=Homo sapiens GN=EEF1D PE=1 SV=5	16	15	2.0	0.014		15429.3	16613.0	17412.8	43274.2	51768.4	67419.1	56048.1	49274.3	63702.9	23185.6	27276.9	31575.3	12336.4	26713.4	19910.6	15420.5	15483.3	14332.5
IF4A1_HUMAN	Eukaryotic initiation factor 4A-I OS=Homo sapiens GN=EIF4A1 PE=1 SV=1	33	17	2.0	0.256		39197.7	40545.3	39690.4	90038.4	82270.4	86967.2	728604.5	756377.7	720456.2	35440.6	39511.6	42384.2	307838.9	390980.4	298220.9	47895.7	52232.5	50144.4
ROA2_HUMAN	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1 SV=2	7	6	2.0	0.119		6333.7	6681.6	6829.9	21382.1	15056.5	8502.8	49796.1	42964.0	56626.7	10394.9	9608.2	8985.4	15227.2	22285.1	19139.9	6586.5	6690.0	6064.1
SAM15_HUMAN	Sterile alpha motif domain-containing protein 15 OS=Homo sapiens GN=SAMD15 PE=2 SV=1	5	1	2.0	0.091		2787.8	2705.8	2832.6	267902.9	261930.2	267514.4	215715.5	227314.4	190044.4	121353.8	128294.9	116385.5	89800.3	118259.0	105435.7	9641.8	9050.9	9499.1
ADT2_HUMAN	ADP/ATP translocase 2 OS=Homo sapiens GN=SLC25A5 PE=1 SV=7	10	2	2.0	0.141		197.3	53.7	294.7	417.9	1022.0	448.8	2292.5	1164.4	1520.5	826.0	320.6	153.8	823.8	91.3	366.3	344.1	320.5	406.9
RS6_HUMAN	40S ribosomal protein S6 OS=Homo sapiens GN=RPS6 PE=1 SV=1	8	7	2.0	0.066		11950.9	12155.9	12018.9	95104.7	120632.0	77735.8	97694.5	94223.5	97050.4	74551.7	60119.4	78242.2	18502.1	21951.2	18725.7	11464.2	10933.1	10692.9
NPM_HUMAN	Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2	7	7	2.0	0.075		4285.8	5015.8	4342.0	18901.0	27521.0	49813.8	35095.2	37346.8	28269.1	25371.5	19406.2	18901.6	7974.5	8823.5	9643.0	4457.3	4899.4	4596.5
SGT1_HUMAN	Protein SGT1 homolog OS=Homo sapiens GN=SUGT1 PE=1 SV=3	4	3	2.0	0.198		852.8	913.8	695.0	11093.9	3241.7	7767.4	4120.9	1906.9	1184.9	3907.8	4000.8	1026.3	1981.7	1402.3	1501.4	524.0	759.5	669.9
EWS_HUMAN	RNA-binding protein EWS OS=Homo sapiens GN=EWSR1 PE=1 SV=1	3	2	2.0	0.090		953.5	1224.1	1161.6	53551.9	55933.5	55510.0	39317.8	43872.0	50251.4	18555.4	25210.2	21911.6	24813.7	29650.1	22133.7	2441.0	2874.1	2577.5
OLA1_HUMAN	Obg-like ATPase 1 OS=Homo sapiens GN=OLA1 PE=1 SV=2	2	1	2.0	0.088		563.4	664.4	646.6	5165.5	10556.7	10696.9	9079.9	9245.0	10385.8	2591.1	2550.8	3601.4	4697.0	6864.0	5855.0	684.1	808.2	738.3
OAT_HUMAN	Ornithine aminotransferase, mitochondrial OS=Homo sapiens GN=OAT PE=1 SV=1	7	7	2.0	0.100		65171.4	64396.3	65232.1	7527096.6	7262394.6	7508387.6	5622238.9	6349160.3	5819711.0	3381616.0	3843854.6	3695450.2	2102251.2	3401947.9	3063985.0	227113.9	258580.1	251642.6
LDHA_HUMAN	L-lactate dehydrogenase A chain OS=Homo sapiens GN=LDHA PE=1 SV=2	27	19	2.0	0.029		21980.0	24050.7	22628.7	104217.3	72799.4	106387.1	91054.9	102065.0	107609.1	72295.6	62716.8	50608.1	29557.3	31078.3	26828.0	17981.5	19444.2	18671.1
TBB3_HUMAN	Tubulin beta-3 chain OS=Homo sapiens GN=TUBB3 PE=1 SV=2	33	2	2.0	0.361		940.4	985.3	927.0	444.5	0.0	0.0	56027.0	61966.3	65473.7	8809.1	14230.4	9011.8	6992.8	28716.2	25196.2	433.2	468.8	542.4
PABP3_HUMAN	Polyadenylate-binding protein 3 OS=Homo sapiens GN=PABPC3 PE=1 SV=2	13	1	2.0	0.072		13974.7	14412.0	14846.0	157960.3	172002.3	162131.7	180425.5	210471.3	210273.3	98708.3	123350.4	94213.6	67902.8	66153.0	69506.9	19019.8	18551.4	18641.1
IF2A_HUMAN	Eukaryotic translation initiation factor 2 subunit 1 OS=Homo sapiens GN=EIF2S1 PE=1 SV=3	3	2	2.0	0.167		2050.4	2348.2	2105.5	106508.6	49887.9	17856.2	78983.7	87365.1	84448.4	55687.5	49521.0	55527.9	13696.6	16791.3	19456.7	2846.2	2668.2	2715.7
EIF3I_HUMAN	Eukaryotic translation initiation factor 3 subunit I OS=Homo sapiens GN=EIF3I PE=1 SV=1	3	3	2.0	0.088		1393.1	1219.5	1443.6	4993.0	5348.6	13277.7	7049.6	5156.3	5538.4	3170.9	3371.6	3986.4	2819.2	3577.1	2125.4	1342.3	1462.1	1204.3
PYGM_HUMAN	Glycogen phosphorylase, muscle form OS=Homo sapiens GN=PYGM PE=1 SV=6	7	1	2.0	0.168		330.7	291.9	194.2	5880.2	13481.0	11257.6	17980.7	18960.9	14772.4	13061.5	13979.2	6160.0	3687.8	977.0	2141.3	771.7	644.4	878.9
IF4H_HUMAN	Eukaryotic translation initiation factor 4H OS=Homo sapiens GN=EIF4H PE=1 SV=5	5	3	2.0	0.089		388.7	570.6	676.5	8493.4	5153.6	4411.5	6079.8	9026.4	7268.1	3748.8	4090.9	3399.1	2547.4	3219.6	2118.9	806.7	738.9	735.6
HNRPR_HUMAN	Heterogeneous nuclear ribonucleoprotein R OS=Homo sapiens GN=HNRNPR PE=1 SV=1	2	1	2.0	0.265		4.7	0.0	72.1	10721.8	12273.2	503.6	5344.6	8862.9	10361.6	5402.6	10090.9	9057.6	0.0	0.0	0.0	12.6	5.4	26.1
F10A1_HUMAN	Hsc70-interacting protein OS=Homo sapiens GN=ST13 PE=1 SV=2	7	1	2.0	0.116		162.5	150.9	151.7	6642.7	10292.1	7743.2	5604.1	5628.6	5429.9	2408.5	3562.7	3006.9	3000.5	4074.9	3892.7	399.0	373.7	648.0
G3P_HUMAN	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	33	30	1.9	0.017		98919.5	97771.3	105341.5	393995.3	167400.7	284312.9	258240.9	252285.6	227223.9	127147.1	160876.5	134721.2	68316.9	104264.5	89548.1	95861.6	91423.2	94919.3
TCPD_HUMAN	T-complex protein 1 subunit delta OS=Homo sapiens GN=CCT4 PE=1 SV=4	26	23	1.9	0.123		19230.1	18115.4	18062.5	775180.2	726685.4	770391.4	673485.5	719085.4	720597.7	586561.3	623156.7	548038.1	143484.8	175060.7	142158.4	20237.3	19388.0	20336.8
PDIA3_HUMAN	Protein disulfide-isomerase A3 OS=Homo sapiens GN=PDIA3 PE=1 SV=4	32	29	1.9	0.119		28657.2	30808.9	29536.9	773201.9	925473.5	932519.8	919873.0	1035465.6	1071686.9	701545.7	735613.2	736968.4	227953.4	249318.3	238538.5	25057.9	25104.7	23467.1
APEX1_HUMAN	DNA-(apurinic or apyrimidinic site) lyase OS=Homo sapiens GN=APEX1 PE=1 SV=2	4	3	1.9	0.156		516.7	622.9	617.2	10229.4	3263.4	2983.0	8486.6	9812.1	3553.3	3458.3	2594.0	2946.2	2286.1	4194.6	2992.2	796.5	756.1	703.4
HNRPQ_HUMAN	Heterogeneous nuclear ribonucleoprotein Q OS=Homo sapiens GN=SYNCRIP PE=1 SV=2	10	8	1.9	0.150		5548.3	5305.1	5332.6	383629.1	440758.8	392966.9	286991.2	348638.6	345347.0	309240.9	321187.9	319613.1	48242.1	60528.5	63900.9	8467.7	8647.6	8485.2
HCN3_HUMAN	otassium/sodium hyperpolarization-activated cyclic nucleotide-gated channel 3 OS=Homo sapiens GN=HCN3 PE=2 SV	2	1	1.9	0.128		25123.3	24111.2	24934.6	78823.1	102706.8	62763.9	182777.9	130214.9	226118.3	153273.7	83942.0	35615.4	34564.9	13336.6	29790.3	31620.9	31891.1	31322.1
RD23B_HUMAN	UV excision repair protein RAD23 homolog B OS=Homo sapiens GN=RAD23B PE=1 SV=1	7	4	1.9	0.102		2881.0	3234.5	2646.9	136709.9	149074.0	144642.2	124045.9	145473.2	145392.8	75715.2	82037.6	79824.0	59162.3	69987.3	59523.4	5723.5	6082.1	5590.0
RCN1_HUMAN	Reticulocalbin-1 OS=Homo sapiens GN=RCN1 PE=1 SV=1	4	2	1.9	0.283		413.1	374.4	478.4	0.0	0.0	469.8	0.0	30.0	178.9	0.0	0.0	0.0	0.0	0.0	0.0	388.5	243.2	385.8
PGK1_HUMAN	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3	24	20	1.9	0.110		29710.2	28407.8	28666.4	851806.9	802246.7	866476.7	532717.0	583161.6	578521.3	390469.5	450245.7	355135.6	253256.2	344871.4	318256.8	47224.6	45894.6	49405.9
UL87_HHV6Z	Protein U58 OS=Human herpesvirus 6B (strain Z29) GN=U58 PE=3 SV=1	3	2	1.9	0.266		1181.4	699.7	370.1	10662.3	354.1	1187.3	9088.8	14875.3	13406.4	7461.4	10086.8	1308.7	1792.5	546.2	3756.2	747.9	737.4	754.6
AT1A2_HUMAN	Sodium/potassium-transporting ATPase subunit alpha-2 OS=Homo sapiens GN=ATP1A2 PE=1 SV=1	13	5	1.9	0.075		5915.4	5519.8	5436.2	41210.9	46032.2	47923.1	35882.1	35587.7	36195.0	29636.8	30200.9	28948.8	9856.0	13648.8	12280.9	3886.0	4200.6	4141.0
ILF2_HUMAN	Interleukin enhancer-binding factor 2 OS=Homo sapiens GN=ILF2 PE=1 SV=2	3	3	1.9	0.221		310.3	85.2	180.1	823.6	377.0	714.7	1605.3	1255.5	2160.0	972.8	1514.2	1121.4	118.5	20.3	0.0	59.9	110.2	74.8
G3BP1_HUMAN	Ras GTPase-activating protein-binding protein 1 OS=Homo sapiens GN=G3BP1 PE=1 SV=1	8	6	1.9	0.168		4023.3	3760.8	4013.9	116289.9	121361.8	108260.5	180259.5	194399.3	184287.9	122923.4	132267.1	133323.6	29378.5	26141.1	31716.2	3811.1	4031.3	4351.1
NASP_HUMAN	Nuclear autoantigenic sperm protein OS=Homo sapiens GN=NASP PE=1 SV=2	25	15	1.9	0.066		156467.2	153809.0	162391.8	934039.2	903773.3	914587.8	547081.8	651726.8	609318.4	544363.8	579581.8	552830.5	218095.7	247555.6	210141.6	111237.1	114862.0	117469.9
TCPZ_HUMAN	T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3	22	15	1.9	0.122		17165.6	18081.6	19002.6	1266845.6	1175724.5	1211678.9	1032924.9	1133537.0	1037109.8	644742.9	801011.0	703542.4	407614.1	524663.7	499595.7	41990.7	41143.9	42893.5
IF2B2_HUMAN	Insulin-like growth factor 2 mRNA-binding protein 2 OS=Homo sapiens GN=IGF2BP2 PE=1 SV=2	5	3	1.9	0.153		761.8	975.7	731.1	27079.0	31569.8	33557.6	36494.8	45978.8	36555.0	31408.5	33741.3	14657.1	9735.9	11566.9	10825.8	1208.0	1021.1	1056.4
ANXA1_HUMAN	Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2	20	13	1.9	0.082		15034.8	15795.3	15265.9	81989.8	115790.5	93873.2	155997.1	75303.7	97943.2	53833.0	35694.4	56056.4	43218.7	66726.8	45891.3	19296.0	19862.3	19765.4
GUAA_HUMAN	GMP synthase [glutamine-hydrolyzing] OS=Homo sapiens GN=GMPS PE=1 SV=1	2	1	1.8	0.181		458.0	1450.6	1385.4	32937.9	22761.3	25419.8	26751.6	32429.6	37044.0	27097.7	30191.0	26330.0	3310.7	2525.1	4977.7	1804.7	860.5	999.1
RSSA_HUMAN	40S ribosomal protein SA OS=Homo sapiens GN=RPSA PE=1 SV=4	12	11	1.8	0.064		25147.9	26684.5	26653.7	78065.7	68278.9	119070.8	145194.1	146052.4	127847.5	83328.0	39140.8	77490.6	52583.5	57773.9	41028.7	21039.7	21228.3	22159.3
SYTC_HUMAN	Threonine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=TARS PE=1 SV=3	15	12	1.8	0.126		8978.1	9131.5	9054.0	827297.5	818159.7	857801.7	745789.6	771606.2	848970.3	434865.6	489464.1	449128.8	339961.7	438899.9	414514.6	37161.9	35813.0	35981.2
LSR_HUMAN	Lipolysis-stimulated lipoprotein receptor OS=Homo sapiens GN=LSR PE=1 SV=4	4	2	1.8	0.037		9453.7	7814.9	8287.5	20248.9	25439.0	3000.7	14960.4	14386.6	21728.0	9588.5	11824.5	5479.8	7647.9	4868.7	4204.5	7818.2	8691.7	8557.1
PIPNA_HUMAN	Phosphatidylinositol transfer protein alpha isoform OS=Homo sapiens GN=PITPNA PE=1 SV=2	2	2	1.8	0.127		382.7	340.3	286.4	10538.0	13445.6	16632.7	14690.0	13946.9	13412.8	6416.1	7075.0	7141.1	7137.0	7627.4	8065.8	782.8	976.3	856.4
RUVB1_HUMAN	RuvB-like 1 OS=Homo sapiens GN=RUVBL1 PE=1 SV=1	9	7	1.8	0.132		5111.5	5276.7	5118.8	287807.4	267783.2	276121.1	295920.1	323371.4	322540.8	186154.5	196930.9	173238.3	116046.2	148438.6	130466.8	11577.6	11808.0	12065.3
UCHL1_HUMAN	Ubiquitin carboxyl-terminal hydrolase isozyme L1 OS=Homo sapiens GN=UCHL1 PE=1 SV=2	6	4	1.8	0.554		76.8	178.9	63.0	3783.2	26681.9	2728.9	1237.2	1602.7	1451.0	8222.6	1486.2	8570.6	1007.1	284.4	1055.1	96.7	111.2	98.8
ALDOA_HUMAN	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2	28	23	1.8	0.018		40984.3	43155.5	42555.4	93692.5	121341.1	123689.1	116810.0	143191.0	139288.0	64453.0	73497.3	66059.7	41463.7	69408.3	42984.5	41931.2	42053.7	39897.3
MYH11_HUMAN	Myosin-11 OS=Homo sapiens GN=MYH11 PE=1 SV=3	14	8	1.8	0.145		99749.0	104150.0	105996.6	10333086.6	10533944.9	11081581.8	7157941.8	8091226.1	8220847.7	4314670.9	4789767.0	4504607.7	4677585.1	5994126.6	5232131.0	521403.2	516339.4	531344.0
CN119_HUMAN	Uncharacterized protein C14orf119 OS=Homo sapiens GN=C14orf119 PE=1 SV=1	1	1	1.8	0.328		1126.7	1162.2	1150.2	3626.8	5561.9	4934.8	32131.6	34119.9	24968.2	1606.5	3481.4	1541.4	15493.9	16638.5	16056.5	2103.2	1873.9	1922.2
RTCB_HUMAN	tRNA-splicing ligase RtcB homolog OS=Homo sapiens GN=RTCB PE=1 SV=1	5	4	1.8	0.137		4641.2	4691.7	4495.8	86111.5	74864.0	69631.0	83099.5	83625.3	90342.8	67669.5	58803.7	64939.0	25180.5	27346.0	21542.7	5326.1	4516.6	4946.8
RAB1C_HUMAN	Putative Ras-related protein Rab-1C OS=Homo sapiens GN=RAB1C PE=5 SV=2	2	1	1.8	0.191		24.9	55.3	47.6	797.3	1625.8	990.4	695.5	2031.8	1516.7	488.8	585.0	503.9	754.4	514.6	1138.2	127.6	136.4	120.4
PRDX6_HUMAN	Peroxiredoxin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3	7	6	1.8	0.143		12363.3	12549.4	12634.4	1168250.5	1079215.2	1185848.8	885832.4	971711.9	903457.6	516890.4	557702.3	516911.7	484909.1	656831.5	606680.9	53986.7	52223.8	54380.4
LMAN1_HUMAN	Protein ERGIC-53 OS=Homo sapiens GN=LMAN1 PE=1 SV=2	5	3	1.8	0.176		1933.6	1758.9	2192.0	38509.0	69517.3	45237.2	53402.4	65924.8	67452.9	53913.7	39793.1	45323.1	14691.2	20184.1	15492.0	1657.4	1921.4	1861.5
NMT1_HUMAN	Glycylpeptide N-tetradecanoyltransferase 1 OS=Homo sapiens GN=NMT1 PE=1 SV=2	2	1	1.8	0.176		141.2	131.8	165.8	4815.8	2075.1	4415.9	4457.6	6623.9	4621.0	2660.5	3596.8	3086.1	1812.8	1725.2	1973.7	131.3	295.5	183.2
PDIA6_HUMAN	Protein disulfide-isomerase A6 OS=Homo sapiens GN=PDIA6 PE=1 SV=1	14	13	1.8	0.335		18199.1	17751.7	15410.0	65733.6	72509.6	78793.8	508655.6	572959.6	516151.1	57776.0	61830.7	51155.8	249870.2	283781.8	242034.4	36188.4	35979.7	37418.3
EIF3D_HUMAN	Eukaryotic translation initiation factor 3 subunit D OS=Homo sapiens GN=EIF3D PE=1 SV=1	3	2	1.8	0.286		314.5	309.1	271.1	17691.2	18315.0	18411.3	32160.6	54992.2	40313.0	20672.2	40035.9	25596.4	7689.5	4864.1	3375.7	502.4	461.3	397.4
6PGD_HUMAN	6-phosphogluconate dehydrogenase, decarboxylating OS=Homo sapiens GN=PGD PE=1 SV=3	11	9	1.8	0.165		7740.4	8025.1	6564.3	213607.7	206191.4	226546.8	317404.4	371034.8	335494.7	111273.0	130928.7	112889.5	149912.9	225514.8	176022.6	17721.6	19153.0	18580.6
NO66_HUMAN	Bifunctional lysine-specific demethylase and histidyl-hydroxylase NO66 OS=Homo sapiens GN=NO66 PE=1 SV=2	4	1	1.8	0.328		1636.9	1729.5	1732.0	7418.2	6623.5	6640.2	54833.2	58738.2	50623.5	16015.6	16125.2	18514.6	17215.4	17043.0	17218.2	1838.5	1949.0	2057.4
ANXA6_HUMAN	Annexin A6 OS=Homo sapiens GN=ANXA6 PE=1 SV=3	6	5	1.8	0.246		166.1	161.8	289.9	17479.9	20245.9	19771.7	28756.4	29913.7	23729.2	19196.3	28381.1	23895.4	2156.2	2605.0	1961.7	609.3	515.4	590.8
RLA0_HUMAN	60S acidic ribosomal protein P0 OS=Homo sapiens GN=RPLP0 PE=1 SV=1	11	3	1.8	0.405		5072.0	4774.9	4504.8	0.0	0.0	0.0	204.6	0.0	1042.6	232.8	615.2	0.0	108.2	356.5	0.2	2611.4	2556.8	2394.1
IF4A3_HUMAN	Eukaryotic initiation factor 4A-III OS=Homo sapiens GN=EIF4A3 PE=1 SV=4	11	7	1.8	0.116		4952.2	5352.8	4916.1	10529.5	11534.7	14759.5	27542.7	28456.8	28793.8	6341.4	10858.1	5355.5	10278.0	18047.6	14730.1	4376.1	3974.0	4188.5
EMD_HUMAN	Emerin OS=Homo sapiens GN=EMD PE=1 SV=1	4	4	1.7	0.093		2361.0	2678.5	2691.5	20496.7	18464.6	24627.2	14062.5	21669.2	21599.5	12633.1	11120.2	10157.9	9416.1	10494.7	9340.4	3133.8	3753.2	3545.2
PAP1L_HUMAN	Polyadenylate-binding protein 1-like OS=Homo sapiens GN=PABPC1L PE=2 SV=1	7	2	1.7	0.184		541.8	326.1	445.6	2012.8	527.9	2680.8	3388.6	4756.8	4906.5	1606.7	1910.7	1940.0	967.1	656.1	177.8	1302.0	1259.0	1393.3
TBB6_HUMAN	Tubulin beta-6 chain OS=Homo sapiens GN=TUBB6 PE=1 SV=1	27	2	1.7	0.411		881.3	938.3	983.3	884.1	592.7	415.4	31189.3	30193.0	27519.2	11525.4	13439.7	14108.4	6175.8	3822.5	2606.5	757.7	671.7	641.5
CISY_HUMAN	Citrate synthase, mitochondrial OS=Homo sapiens GN=CS PE=1 SV=2	18	14	1.7	0.135		18309.8	18517.8	21172.7	125175.2	89063.8	149623.0	126215.4	216612.5	176147.1	47632.3	75616.9	46302.4	108882.5	95613.4	115409.7	16954.9	17394.4	17401.9
EIF2A_HUMAN	Eukaryotic translation initiation factor 2A OS=Homo sapiens GN=EIF2A PE=1 SV=3	1	1	1.7	0.319		44.8	24.9	7.2	23529.2	14395.7	26046.9	29560.3	30567.4	147.2	22970.3	18740.3	20899.8	5104.0	0.0	3890.8	22.8	0.3	39.3
API5_HUMAN	Apoptosis inhibitor 5 OS=Homo sapiens GN=API5 PE=1 SV=3	4	2	1.7	0.217		967.6	1056.9	1735.8	47560.6	22607.8	51753.9	53513.8	57782.3	55143.6	38407.3	50212.3	40899.0	9441.1	13918.1	12099.2	1364.6	1120.0	1138.8
SERA_HUMAN	D-3-phosphoglycerate dehydrogenase OS=Homo sapiens GN=PHGDH PE=1 SV=4	13	12	1.7	0.197		6388.6	6616.9	7145.9	259280.9	232179.9	232996.4	240615.1	281144.2	308104.4	217968.9	244816.8	235922.1	57098.4	63285.8	71346.1	6845.7	6040.0	5893.9
CAP1_HUMAN	Adenylyl cyclase-associated protein 1 OS=Homo sapiens GN=CAP1 PE=1 SV=5	10	8	1.7	0.155		5876.2	5862.7	6037.1	301904.4	311266.0	313943.8	250655.9	284678.9	288409.9	191911.7	167045.1	176955.9	143031.8	164488.0	136800.2	16271.2	14996.9	15409.5
IMDH2_HUMAN	Inosine-5'-monophosphate dehydrogenase 2 OS=Homo sapiens GN=IMPDH2 PE=1 SV=2	12	10	1.7	0.204		5530.0	5673.3	5402.5	115149.5	137785.2	127197.0	150653.4	170150.5	148434.2	119606.5	137894.4	139629.8	31036.2	33706.1	28559.6	4347.8	4121.8	4336.6
RAB1B_HUMAN	Ras-related protein Rab-1B OS=Homo sapiens GN=RAB1B PE=1 SV=1	2	1	1.7	0.356		18.0	4.5	5.9	924.5	909.9	47.5	731.2	445.6	584.1	553.0	782.9	762.1	38.4	0.0	0.0	1.9	15.2	5.5
ACTBM_HUMAN	Putative beta-actin-like protein 3 OS=Homo sapiens GN=POTEKP PE=5 SV=1	10	2	1.7	0.173		82678.0	83805.7	84446.8	11676805.0	11136762.5	11498197.4	8463239.8	9444272.1	10335963.0	5059735.3	5589157.9	5247027.0	5434971.6	7285557.4	6628397.2	567987.3	571623.9	588359.0
RINI_HUMAN	Ribonuclease inhibitor OS=Homo sapiens GN=RNH1 PE=1 SV=2	3	2	1.7	0.196		505.4	481.9	523.4	0.0	432.5	0.0	1180.7	317.4	947.7	0.0	0.0	265.3	151.0	244.3	520.1	515.0	453.5	437.5
PSD13_HUMAN	26S proteasome non-ATPase regulatory subunit 13 OS=Homo sapiens GN=PSMD13 PE=1 SV=2	4	3	1.7	0.347		95.6	154.5	190.6	3407.3	860.3	1037.5	635.6	412.0	2564.8	1245.4	1100.7	374.8	1350.7	758.0	581.0	14.7	64.1	96.2
GDIB_HUMAN	Rab GDP dissociation inhibitor beta OS=Homo sapiens GN=GDI2 PE=1 SV=2	11	6	1.7	0.239		10129.4	10290.4	10329.0	846525.4	773784.5	773622.0	689183.4	747973.0	672233.0	710158.7	788961.1	688200.8	131788.6	164376.2	148148.8	27395.8	26443.0	26871.3
ILF3_HUMAN	Interleukin enhancer-binding factor 3 OS=Homo sapiens GN=ILF3 PE=1 SV=3	6	6	1.7	0.260		1870.6	2039.8	1770.0	73753.0	202369.4	79398.1	117098.1	108346.6	86983.1	90029.1	97609.8	85717.7	39830.4	44440.5	32009.0	4582.0	4517.4	4304.6
PSMD4_HUMAN	26S proteasome non-ATPase regulatory subunit 4 OS=Homo sapiens GN=PSMD4 PE=1 SV=1	6	6	1.7	0.090		15509.5	17815.2	17009.4	60542.7	79771.1	90157.0	91422.1	112135.1	108343.6	38827.1	43344.0	43356.3	55144.6	58111.9	47645.0	22419.2	23844.5	22751.1
SMC1A_HUMAN	Structural maintenance of chromosomes protein 1A OS=Homo sapiens GN=SMC1A PE=1 SV=2	3	3	1.7	0.227		1497.6	1572.1	1564.4	80384.2	125464.8	121079.8	63532.8	73546.9	75637.4	24762.0	29986.2	32770.8	68039.0	84744.9	66217.1	7168.8	6617.3	6485.9
RL3_HUMAN	60S ribosomal protein L3 OS=Homo sapiens GN=RPL3 PE=1 SV=2	15	13	1.7	0.178		9964.7	10789.9	10439.6	152463.4	137857.3	150717.4	169373.5	204804.2	193005.0	60338.7	61686.8	62934.1	145111.8	103486.4	146462.6	15116.5	16289.5	15656.4
HNRH2_HUMAN	Heterogeneous nuclear ribonucleoprotein H2 OS=Homo sapiens GN=HNRNPH2 PE=1 SV=1	6	3	1.7	0.394		1662.1	2026.6	1823.1	4778.3	7179.8	4379.9	63055.6	61079.0	60867.6	16454.7	26405.6	22122.7	19261.1	17719.2	16295.6	2588.7	2157.4	2174.0
TADBP_HUMAN	TAR DNA-binding protein 43 OS=Homo sapiens GN=TARDBP PE=1 SV=1	4	4	1.6	0.073		1785.1	1620.8	1625.0	2416.8	5314.2	6119.9	5080.8	5929.1	4440.8	1202.4	776.0	2017.6	4162.4	4482.0	3083.5	1651.4	1889.7	1566.0
DDX6_HUMAN	Probable ATP-dependent RNA helicase DDX6 OS=Homo sapiens GN=DDX6 PE=1 SV=2	3	3	1.6	0.282		680.2	649.3	694.9	13335.7	24441.1	10939.3	26199.1	24285.8	28301.8	20322.1	19937.6	23343.3	4013.4	4756.1	4253.7	834.7	646.7	797.6
SSRP1_HUMAN	FACT complex subunit SSRP1 OS=Homo sapiens GN=SSRP1 PE=1 SV=1	9	7	1.6	0.197		4074.4	4856.1	4032.3	243935.0	288933.6	291445.5	203328.9	231297.4	245456.1	105622.4	128855.8	122337.9	154790.9	191494.2	177900.4	14720.6	15762.5	14541.2
TAGL2_HUMAN	Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3	3	2	1.6	0.302		766.9	800.7	732.1	2577.7	61131.0	75066.8	37738.7	48954.9	48747.5	28606.0	30382.1	33166.3	20298.5	27668.9	24811.4	2082.0	1696.7	2005.7
SYHC_HUMAN	Histidine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=HARS PE=1 SV=2	4	1	1.6	0.590		379.9	386.8	313.5	0.0	0.0	0.0	2845.1	6095.1	6858.6	8529.2	840.4	171.2	0.0	141.5	0.0	334.2	206.5	199.0
PDIA1_HUMAN	Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3	33	29	1.6	0.211		23083.4	23809.3	23558.1	1206807.3	1218057.7	1249360.3	978105.1	1246826.8	1110700.9	840716.6	970772.3	763243.1	493138.6	603443.8	574492.4	49273.3	50078.0	50515.6
TCPB_HUMAN	T-complex protein 1 subunit beta OS=Homo sapiens GN=CCT2 PE=1 SV=4	33	27	1.6	0.220		46261.4	44317.8	47947.6	773793.9	673166.7	697995.3	750610.5	775157.4	742615.7	684608.9	758285.1	591308.2	202871.3	208156.1	220048.5	54771.7	56381.1	57286.5
NACAM_HUMAN	Nascent polypeptide-associated complex subunit alpha, muscle-specific form OS=Homo sapiens GN=NACA PE=1 SV=1	19	11	1.6	0.203		8042.6	9076.1	8836.6	252871.9	221712.3	200326.8	172404.5	182601.9	163062.7	84916.4	104444.3	76279.7	130347.7	161877.0	147646.7	17784.0	18881.4	18465.7
PA2G4_HUMAN	Proliferation-associated protein 2G4 OS=Homo sapiens GN=PA2G4 PE=1 SV=3	20	17	1.6	0.280		24250.9	25517.2	25031.0	119377.1	171283.6	215201.8	327118.5	418477.5	385393.7	74774.1	70218.5	76547.5	222810.8	278558.5	260805.2	29776.2	32801.7	31309.0
PAK2_HUMAN	Serine/threonine-protein kinase PAK 2 OS=Homo sapiens GN=PAK2 PE=1 SV=3	6	4	1.6	0.224		13823.7	14485.7	15144.6	2765242.9	2498017.5	2553536.1	2005226.3	2115132.5	2358498.4	1219970.5	1302518.8	1227312.9	1302008.4	1913493.0	1628329.1	140591.3	155320.6	147171.3
PRS4_HUMAN	26S protease regulatory subunit 4 OS=Homo sapiens GN=PSMC1 PE=1 SV=1	7	6	1.6	0.271		6827.7	6879.7	6803.8	83340.7	107295.8	123095.7	62048.1	53644.2	73999.4	22168.5	30661.3	65594.9	111633.3	44148.0	33566.0	8925.0	9111.5	7924.4
DDX1_HUMAN	ATP-dependent RNA helicase DDX1 OS=Homo sapiens GN=DDX1 PE=1 SV=2	4	2	1.6	0.514		6.3	0.0	0.0	5049.3	4223.0	2229.0	1803.9	0.0	1245.0	2338.1	1652.2	5227.4	63.4	0.0	0.0	0.0	0.0	6.1
ENOG_HUMAN	Gamma-enolase OS=Homo sapiens GN=ENO2 PE=1 SV=3	8	4	1.6	0.277		3114.9	3032.8	2798.0	49974.4	55698.0	80228.9	132113.4	104946.6	104142.6	36262.2	45500.3	51415.2	47307.6	87275.0	55813.4	6469.7	6933.7	6501.0
WDR1_HUMAN	WD repeat-containing protein 1 OS=Homo sapiens GN=WDR1 PE=1 SV=4	6	3	1.6	0.370		1021.4	836.2	1078.1	57435.1	31958.7	33612.1	57266.0	87521.3	78636.9	61960.0	65622.2	66015.0	9774.7	9512.6	9074.2	988.3	965.7	1065.1
UBP7_HUMAN	Ubiquitin carboxyl-terminal hydrolase 7 OS=Homo sapiens GN=USP7 PE=1 SV=2	2	2	1.5	0.261		1967.3	1987.4	2322.3	172006.3	151996.7	181665.7	126702.0	139761.2	152667.4	67002.5	76408.3	64890.2	105511.0	143544.1	124876.9	6876.5	6681.7	6522.5
LDHB_HUMAN	L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2	28	24	1.5	0.044		40567.6	41965.6	39879.2	96249.2	117193.5	113946.5	103909.8	122008.1	131655.9	54460.1	67287.6	61792.9	65753.3	75720.2	77093.0	41795.4	42038.7	41682.3
C1QBP_HUMAN	Complement component 1 Q subcomponent-binding protein, mitochondrial OS=Homo sapiens GN=C1QBP PE=1 SV=1	4	4	1.5	0.542		2720.0	2582.7	2596.4	0.0	0.0	0.0	0.0	0.0	328.1	261.7	0.0	0.0	125.7	18.9	192.5	1681.1	1564.6	1565.9
THIL_HUMAN	Acetyl-CoA acetyltransferase, mitochondrial OS=Homo sapiens GN=ACAT1 PE=1 SV=1	10	9	1.5	0.157		7551.0	7382.9	7228.4	6583.2	15547.7	24915.9	8761.9	9578.9	24828.6	4180.5	4142.4	2896.2	9694.7	15266.9	7678.7	10636.0	9611.2	9898.7
G6PI_HUMAN	Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4	21	18	1.5	0.354		18708.0	18494.7	19497.7	564069.9	592514.0	376689.5	721776.0	598089.7	967321.5	461731.1	837096.5	685453.5	144245.6	214762.9	152380.1	20744.3	19348.5	18632.9
CL054_HUMAN	Uncharacterized protein C12orf54 OS=Homo sapiens GN=C12orf54 PE=2 SV=2	1	1	1.5	0.454		4971.1	4908.3	5363.7	84341.2	77809.0	57114.1	241242.0	260926.0	268761.9	57607.9	260508.3	217780.4	33256.5	41136.6	39452.9	4711.5	4198.2	4292.0
CLIC1_HUMAN	Chloride intracellular channel protein 1 OS=Homo sapiens GN=CLIC1 PE=1 SV=4	7	5	1.5	0.303		339.1	492.8	407.6	3176.5	2659.6	5860.6	8028.7	8208.8	10247.3	2427.1	4611.7	3939.7	4111.4	4159.4	3482.1	1254.7	1229.0	935.8
AT1A3_HUMAN	Sodium/potassium-transporting ATPase subunit alpha-3 OS=Homo sapiens GN=ATP1A3 PE=1 SV=3	9	1	1.5	0.422		2829.9	2853.4	2951.7	75421.5	73123.1	75629.7	150567.3	186455.8	183030.6	143667.1	156490.5	135869.1	18567.0	21744.5	21097.7	1997.6	1973.9	2150.2
HMCS1_HUMAN	Hydroxymethylglutaryl-CoA synthase, cytoplasmic OS=Homo sapiens GN=HMGCS1 PE=1 SV=2	6	4	1.5	0.359		3066.3	3284.5	3218.9	51407.1	45568.8	43901.7	80033.8	91492.6	80078.5	70774.8	79462.8	58129.5	16575.5	19290.7	18776.2	2027.6	1925.4	2134.1
EIF3F_HUMAN	Eukaryotic translation initiation factor 3 subunit F OS=Homo sapiens GN=EIF3F PE=1 SV=1	2	1	1.5	0.303		88.6	376.1	343.6	4279.4	2828.8	2567.7	5886.2	6038.7	4874.1	1056.9	2257.6	1789.0	3062.4	4624.6	3657.3	636.4	588.0	609.9
LRC59_HUMAN	Leucine-rich repeat-containing protein 59 OS=Homo sapiens GN=LRRC59 PE=1 SV=1	4	3	1.5	0.398		473.8	423.7	601.7	5834.8	3689.2	2743.3	14960.2	5390.2	6859.3	4215.1	2765.9	2938.5	4401.2	5586.3	6164.6	547.0	375.3	540.8
NDE1_HUMAN	Nuclear distribution protein nudE homolog 1 OS=Homo sapiens GN=NDE1 PE=1 SV=2	3	2	1.5	0.305		766.1	950.1	756.3	17699.4	17292.9	18217.8	21504.3	23866.9	26414.0	9139.6	10753.9	10764.8	23381.0	5840.6	19559.6	2236.8	2511.0	1998.9
TBB5_HUMAN	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2	62	7	1.5	0.524		66204.1	64510.0	67139.4	93391.4	91438.8	88581.5	2134430.4	2329821.9	2590991.1	1080865.9	1178642.3	1066898.2	488763.8	578547.1	535196.2	54053.2	55117.6	55811.2
VATB2_HUMAN	V-type proton ATPase subunit B, brain isoform OS=Homo sapiens GN=ATP6V1B2 PE=1 SV=3	8	7	1.5	0.374		6617.6	6625.4	6523.3	88159.2	36168.8	25788.0	103915.6	107598.0	128332.0	89825.6	87132.0	60079.4	33409.9	27074.9	26387.3	7110.5	6829.3	7156.3
KPYR_HUMAN	Pyruvate kinase PKLR OS=Homo sapiens GN=PKLR PE=1 SV=2	8	5	1.5	0.300		2203.7	2270.9	2243.9	547649.2	476719.2	508625.7	418636.5	412207.9	535216.8	251586.5	250833.1	218919.9	321493.2	437901.6	381810.8	35404.8	37957.1	36522.8
EF1G_HUMAN	Elongation factor 1-gamma OS=Homo sapiens GN=EEF1G PE=1 SV=3	28	27	1.5	0.257		181555.6	196981.0	233275.7	1056949.2	833533.5	1108096.2	1828272.2	2026569.1	1317418.1	1075668.4	684877.2	934773.7	718604.8	1064732.4	921880.7	195344.5	184024.7	213105.2
NP1L1_HUMAN	Nucleosome assembly protein 1-like 1 OS=Homo sapiens GN=NAP1L1 PE=1 SV=1	8	6	1.5	0.402		5991.6	6071.5	5867.6	62413.8	57673.5	70637.6	165329.9	177459.3	177247.9	106761.1	131361.5	106302.9	50555.5	46531.3	43803.9	4726.9	4726.4	4844.5
HS905_HUMAN	Putative heat shock protein HSP 90-alpha A5 OS=Homo sapiens GN=HSP90AA5P PE=2 SV=1	17	1	1.5	0.634		556.3	595.4	468.6	0.0	0.0	0.0	7349.4	24525.7	25563.1	5258.7	12228.4	18842.9	272.0	2185.3	1720.1	14.5	50.2	0.0
PLSI_HUMAN	Plastin-1 OS=Homo sapiens GN=PLS1 PE=1 SV=2	14	4	1.5	0.205		8530.0	10014.4	10178.0	25197.6	35291.9	22347.7	42155.0	42410.0	50386.2	17925.5	16659.3	15149.3	38494.1	5803.5	35364.7	13793.2	13173.1	13612.7
PSMD3_HUMAN	26S proteasome non-ATPase regulatory subunit 3 OS=Homo sapiens GN=PSMD3 PE=1 SV=2	10	8	1.4	0.461		1849.9	1763.9	1882.3	46176.8	37723.4	24300.1	70712.7	115758.5	87291.4	67469.0	84834.4	65088.2	17092.1	12545.3	15090.9	2338.5	1692.6	1766.6
RPR1B_HUMAN	Regulation of nuclear pre-mRNA domain-containing protein 1B OS=Homo sapiens GN=RPRD1B PE=1 SV=1	2	2	1.4	0.506		136.7	106.2	174.4	202.6	0.0	0.0	801.8	363.9	29.8	0.0	147.0	365.0	74.1	121.9	113.4	151.6	138.6	144.2
GSHR_HUMAN	Glutathione reductase, mitochondrial OS=Homo sapiens GN=GSR PE=1 SV=2	5	5	1.4	0.501		2073.6	1973.2	2012.4	21769.2	8281.6	7562.6	59353.0	63473.2	45082.6	27950.2	57030.2	22287.6	14855.8	7873.0	10603.8	2421.4	1874.6	2226.8
VDAC2_HUMAN	Voltage-dependent anion-selective channel protein 2 OS=Homo sapiens GN=VDAC2 PE=1 SV=2	8	6	1.4	0.169		6262.8	8509.0	8687.9	17873.3	30699.9	34385.6	10126.1	21973.3	25645.4	15616.0	14533.4	15494.0	6639.4	5140.0	14027.7	14237.4	14446.2	14311.8
PTBP1_HUMAN	Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1 PE=1 SV=1	9	9	1.4	0.434		3787.2	4427.4	3972.6	74434.1	75081.6	66588.3	156052.6	178734.4	191293.9	135459.5	135087.6	125829.0	37933.5	32149.8	43315.4	5566.3	5359.2	5519.5
LS14B_HUMAN	Protein LSM14 homolog B OS=Homo sapiens GN=LSM14B PE=1 SV=1	1	1	1.4	0.669		149.0	125.3	109.2	4939.8	2055.9	519.2	2273.6	2370.9	1174.8	8132.0	247.9	0.0	95.2	605.2	71.9	152.8	173.6	184.3
IF4A2_HUMAN	Eukaryotic initiation factor 4A-II OS=Homo sapiens GN=EIF4A2 PE=1 SV=2	17	1	1.4	0.369		392.1	224.4	182.9	9771.3	18280.7	19630.0	11022.0	12794.2	13484.7	6039.1	3824.6	5228.0	12860.0	14341.5	11905.8	2177.6	2099.1	1957.2
PUR9_HUMAN	Bifunctional purine biosynthesis protein PURH OS=Homo sapiens GN=ATIC PE=1 SV=3	14	11	1.4	0.333		21820.7	22084.6	23773.6	245669.3	287887.2	254155.8	380312.8	413599.9	399304.0	190202.2	202860.9	156317.5	238402.2	312400.8	264732.8	30733.0	30145.0	29060.1
ERF1_HUMAN	Eukaryotic peptide chain release factor subunit 1 OS=Homo sapiens GN=ETF1 PE=1 SV=3	2	2	1.4	0.578		1147.1	1291.2	1473.1	171.0	217.2	347.5	39035.8	36371.2	34100.4	14723.9	16239.8	15612.2	9767.1	12032.4	9438.8	1233.8	1040.5	1237.5
EIF3L_HUMAN	Eukaryotic translation initiation factor 3 subunit L OS=Homo sapiens GN=EIF3L PE=1 SV=1	1	1	1.4	0.532		63.6	0.0	12.6	14893.4	2288.1	11783.1	29073.9	21329.8	24026.4	18450.7	25945.0	20793.3	4063.9	1606.8	1984.1	100.7	360.3	465.9
1433Z_HUMAN	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1	10	7	1.4	0.440		618.9	726.7	552.7	19794.8	20736.5	45275.9	25623.9	15447.6	14806.7	17666.7	22880.2	19619.2	6887.5	18795.3	13841.0	1086.5	945.2	1004.4
BASP1_HUMAN	Brain acid soluble protein 1 OS=Homo sapiens GN=BASP1 PE=1 SV=2	8	8	1.4	0.458		8662.2	9874.2	8963.7	83153.2	100420.1	109683.7	254176.0	268603.0	272473.1	160627.8	198559.6	185050.6	70420.0	88624.2	78225.0	7767.3	7717.6	7293.6
ANX11_HUMAN	Annexin A11 OS=Homo sapiens GN=ANXA11 PE=1 SV=1	7	7	1.4	0.446		3805.2	3985.6	3985.2	10779.8	18652.9	20934.8	34015.4	49543.1	53333.1	33718.0	36309.6	28581.0	11086.4	14577.7	9771.9	3365.8	3249.1	2950.5
CAZA1_HUMAN	F-actin-capping protein subunit alpha-1 OS=Homo sapiens GN=CAPZA1 PE=1 SV=3	3	2	1.4	0.315		2053.2	2327.2	2178.5	5770.8	5714.2	4545.3	3198.3	7063.1	9629.9	1655.7	7749.2	8561.6	2679.1	3012.1	2027.8	1794.1	1634.1	1658.9
RAN_HUMAN	GTP-binding nuclear protein Ran OS=Homo sapiens GN=RAN PE=1 SV=3	2	1	1.4	0.614		0.0	0.0	0.0	170.9	554.0	466.1	379.6	1191.1	190.8	681.9	952.0	0.0	12.8	368.5	126.4	0.0	1.4	0.0
TCPA_HUMAN	T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1	22	17	1.4	0.445		8388.7	8022.3	8389.7	371311.3	249835.6	301265.7	367813.8	448406.7	452625.0	380436.7	445796.8	381290.1	108475.1	130132.5	117585.7	15663.9	14875.4	15921.0
CKAP4_HUMAN	Cytoskeleton-associated protein 4 OS=Homo sapiens GN=CKAP4 PE=1 SV=2	18	10	1.4	0.347		2724.9	16658.6	17016.1	252571.5	241463.4	234891.5	217849.7	256821.3	265278.6	124412.5	145422.5	169517.9	167978.5	222492.0	191399.0	24463.8	25833.1	23514.8
HNRPK_HUMAN	Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK PE=1 SV=1	24	21	1.4	0.468		14606.2	13222.7	12840.1	319392.9	404996.9	355706.5	725240.2	818952.6	693834.8	517278.5	619077.7	580448.6	191033.0	249584.3	240322.2	20800.8	21181.7	21493.7
4F2_HUMAN	4F2 cell-surface antigen heavy chain OS=Homo sapiens GN=SLC3A2 PE=1 SV=3	4	3	1.3	0.423		8050.9	8015.0	8455.9	545372.4	532566.9	581617.8	613112.2	633255.8	691713.5	267220.8	299642.5	265602.7	572164.0	613233.2	521923.5	50637.9	49855.9	54701.3
ETFA_HUMAN	Electron transfer flavoprotein subunit alpha, mitochondrial OS=Homo sapiens GN=ETFA PE=1 SV=1	4	4	1.3	0.659		638.8	553.6	614.5	0.0	0.0	0.0	146.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	176.4	381.8	445.0	462.9
AATM_HUMAN	Aspartate aminotransferase, mitochondrial OS=Homo sapiens GN=GOT2 PE=1 SV=3	7	5	1.3	0.251		4387.7	4540.7	4270.4	5148.7	10745.3	10929.2	2169.4	4889.9	5857.0	2821.4	2522.1	3607.2	6954.0	2029.9	8603.5	4607.8	4214.5	4416.3
SAHH_HUMAN	Adenosylhomocysteinase OS=Homo sapiens GN=AHCY PE=1 SV=4	17	15	1.3	0.384		26329.5	24939.6	27303.1	175711.1	252012.6	256831.3	265007.2	184426.8	309552.9	133428.9	100280.2	157230.9	199053.8	232422.2	174315.0	56592.8	56344.0	54381.9
TALDO_HUMAN	Transaldolase OS=Homo sapiens GN=TALDO1 PE=1 SV=2	17	14	1.3	0.289		30733.6	30393.6	31901.1	51616.1	37759.0	52057.2	67439.6	78682.3	80473.9	75874.9	72870.9	69807.0	22776.5	27969.0	28264.6	18502.5	18736.4	19049.8
TPM4_HUMAN	Tropomyosin alpha-4 chain OS=Homo sapiens GN=TPM4 PE=1 SV=3	3	2	1.3	0.451		12468.6	11178.8	13499.5	39387.5	18719.5	28841.4	10401.7	10296.2	7845.4	2272.4	22276.9	33613.7	7693.3	22480.1	9254.9	6687.5	7295.7	6759.0
M3K14_HUMAN	Mitogen-activated protein kinase kinase kinase 14 OS=Homo sapiens GN=MAP3K14 PE=1 SV=2	6	5	1.3	0.336		179169.7	176037.5	176925.7	1070098.0	1144020.3	1046764.1	931475.0	1034714.2	925427.5	826235.7	927481.8	805822.9	508846.3	660276.1	592946.8	293584.4	283422.0	297575.1
PDCD4_HUMAN	Programmed cell death protein 4 OS=Homo sapiens GN=PDCD4 PE=1 SV=2	3	3	1.3	0.683		2466.4	2958.3	3544.5	40433.8	29231.4	16046.7	56279.2	140917.2	153502.6	35035.4	125679.0	126977.4	17639.0	18083.3	14244.4	3832.5	3663.2	3692.0
COPD_HUMAN	Coatomer subunit delta OS=Homo sapiens GN=ARCN1 PE=1 SV=1	1	1	1.3	0.613		371.5	426.3	331.0	7594.3	3113.9	5502.7	14007.7	14171.8	18082.6	6811.8	14162.0	12143.9	4965.2	5072.2	5499.2	703.5	598.9	607.8

GLYM_HUMAN	Serine hydroxymethyltransferase, mitochondrial OS=Homo sapiens GN=SHMT2 PE=1 SV=3	12	8	1.3	0.630		8145.2	8531.1	8540.6	68797.9	105557.3	101369.1	253441.4	287980.9	241322.4	213163.4	231192.7	221907.6	55127.5	65105.5	47923.2	10274.1	10096.0	9818.4
SFXN1_HUMAN	Sideroflexin-1 OS=Homo sapiens GN=SFXN1 PE=1 SV=4	2	2	1.2	0.551		438.6	714.8	487.5	7727.0	6820.4	9507.2	8648.6	6620.3	7910.6	4643.0	2372.2	2275.9	9637.0	8582.9	8768.5	1175.6	939.7	874.2
1433E_HUMAN	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1	17	14	1.2	0.611		4954.4	4510.0	4455.6	51034.7	90085.8	52107.7	71828.6	152273.3	145566.7	101982.4	111704.1	106802.3	38860.1	37266.1	34454.2	10944.9	12317.3	11155.1
PSA1_HUMAN	Proteasome subunit alpha type-1 OS=Homo sapiens GN=PSMA1 PE=1 SV=1	6	6	1.2	0.567		334.3	227.6	257.1	5968.5	9432.0	1658.4	4371.7	7010.8	3342.4	4371.6	4389.8	2949.7	1821.3	505.1	3245.1	3031.0	3127.3	2877.2
HNRPC_HUMAN	Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC PE=1 SV=4	3	2	1.2	0.583		253.6	184.5	203.5	12033.1	5737.6	7578.4	5929.5	8938.8	10662.5	3390.9	6137.9	4617.8	7303.9	10381.9	7582.5	1017.6	635.8	742.7
RPAP3_HUMAN	RNA polymerase II-associated protein 3 OS=Homo sapiens GN=RPAP3 PE=1 SV=2	3	3	1.2	0.444		5928.2	6662.6	6194.3	50101.9	36597.1	30542.1	42029.1	46747.4	33017.7	25271.7	27808.6	32145.9	26248.8	31627.9	34223.1	10383.2	10543.2	11142.3
PSPC1_HUMAN	Paraspeckle component 1 OS=Homo sapiens GN=PSPC1 PE=1 SV=1	3	2	1.2	0.729		217.7	230.9	285.7	5162.9	4596.6	4446.3	29788.9	32666.9	27182.3	21852.2	25153.5	27375.4	3324.4	3099.3	3246.2	412.4	421.6	498.1
PUR1_HUMAN	Amidophosphoribosyltransferase OS=Homo sapiens GN=PPAT PE=1 SV=1	2	2	1.2	0.641		1159.9	1251.4	1422.9	6936.4	8929.7	14017.7	30637.5	37668.5	27811.8	22263.7	23634.4	15735.5	12226.8	13363.0	13024.9	2075.3	1905.5	2147.3
MCM2_HUMAN	DNA replication licensing factor MCM2 OS=Homo sapiens GN=MCM2 PE=1 SV=4	6	2	1.2	0.791		285.8	117.5	326.7	31167.8	8389.9	850.9	22737.1	10920.7	5411.3	10323.2	21222.4	31107.0	988.9	2936.0	182.7	224.9	145.9	193.8
TBB4B_HUMAN	Tubulin beta-4B chain OS=Homo sapiens GN=TUBB4B PE=1 SV=1	64	2	1.2	0.758		24513.0	24570.4	25012.9	54372.6	71937.3	72475.9	682658.7	822032.1	869763.6	479965.7	550178.2	513208.9	183568.5	230732.1	213477.8	20925.7	22473.0	22196.8
AL7A1_HUMAN	Alpha-aminoadipic semialdehyde dehydrogenase OS=Homo sapiens GN=ALDH7A1 PE=1 SV=5	7	5	1.2	0.739		3514.2	3127.0	3651.1	19227.2	26382.7	21132.8	137278.1	145893.4	139591.4	83334.8	101387.5	86762.6	42448.8	51746.2	45520.3	4033.9	3823.5	4141.8
MDHM_HUMAN	Malate dehydrogenase, mitochondrial OS=Homo sapiens GN=MDH2 PE=1 SV=3	15	12	1.2	0.414		17802.0	17281.6	16644.3	59640.7	44113.7	44522.9	37896.6	44553.9	49161.4	27852.0	52811.9	41392.5	27323.5	36750.8	27598.1	23406.9	22702.2	24404.4
HMGB1_HUMAN	High mobility group protein B1 OS=Homo sapiens GN=HMGB1 PE=1 SV=3	9	6	1.2	0.694		282.2	298.0	344.9	5662.5	6909.8	4285.2	6677.0	13253.0	5411.3	5618.9	4512.1	6804.2	5382.9	5784.3	7098.0	719.1	631.7	526.6
1433F_HUMAN	14-3-3 protein eta OS=Homo sapiens GN=YWHAH PE=1 SV=4	6	2	1.2	0.771		85.6	79.5	59.1	850.9	1227.4	765.0	1942.1	3981.4	3887.8	2022.5	3004.8	3033.7	928.1	976.8	858.4	116.0	130.4	79.5
GSTP1_HUMAN	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2	6	6	1.1	0.819		39.0	48.6	79.2	785.2	229.8	68.0	1773.5	1178.8	927.9	696.1	0.0	183.6	311.2	2010.3	867.2	110.1	130.8	198.9
RL7_HUMAN	60S ribosomal protein L7 OS=Homo sapiens GN=RPL7 PE=1 SV=1	12	9	1.1	0.715		4987.1	4870.3	4583.1	38470.9	35737.2	27632.7	61499.5	79415.9	77478.5	45965.8	52728.2	54175.5	37916.7	34598.0	34187.1	11447.6	12601.6	12297.8
AMPL_HUMAN	Cytosol aminopeptidase OS=Homo sapiens GN=LAP3 PE=1 SV=3	5	4	1.1	0.859		1053.0	996.4	1137.2	3772.1	3058.9	1781.3	41918.5	16635.2	31331.5	27345.1	13642.1	26887.7	9851.5	6167.7	4403.9	1132.1	1006.5	1094.7
2ABA_HUMAN	e/threonine-protein phosphatase 2A 55 kDa regulatory subunit B alpha isoform OS=Homo sapiens GN=PPP2R2A PE=1	4	2	1.1	0.829		656.3	755.3	690.4	5353.5	4009.3	5983.3	16850.7	19983.5	21161.1	16346.9	18705.4	16515.5	4130.0	5398.6	3884.9	974.3	893.9	1145.4
MARCS_HUMAN	Myristoylated alanine-rich C-kinase substrate OS=Homo sapiens GN=MARCKS PE=1 SV=4	3	2	1.1	0.868		145.6	125.8	112.9	7760.1	7324.1	4508.1	17375.1	17733.4	16556.8	18514.6	25069.8	20726.9	131.8	0.0	0.0	132.9	270.2	138.4
SERPH_HUMAN	Serpin H1 OS=Homo sapiens GN=SERPINH1 PE=1 SV=2	15	12	1.1	0.855		48280.7	48104.0	51173.3	119812.0	98170.7	107960.4	834923.3	876631.1	1007814.2	136787.0	140241.7	155654.2	625006.2	847686.0	720877.8	91873.9	90833.9	93889.7
DPOD1_HUMAN	DNA polymerase delta catalytic subunit OS=Homo sapiens GN=POLD1 PE=1 SV=2	1	1	1.1	0.860		151.0	170.8	201.6	13445.8	8765.4	5810.0	5835.9	6678.2	2273.8	2527.4	4446.8	4052.5	6255.0	8698.7	12701.0	463.1	403.7	458.2
FSCN1_HUMAN	Fascin OS=Homo sapiens GN=FSCN1 PE=1 SV=3	11	10	1.0	0.933		7446.9	7114.3	8390.0	52711.6	25547.3	28731.3	205790.5	223205.7	186198.2	148669.7	184369.4	172280.2	55578.8	71437.0	60707.4	7043.6	7614.6	7175.5
RANG_HUMAN	Ran-specific GTPase-activating protein OS=Homo sapiens GN=RANBP1 PE=1 SV=1	2	2	1.0	0.860		30154.4	29976.2	33018.8	26101.1	24474.8	18855.3	23636.5	64205.6	55818.9	12306.3	36626.9	52757.3	23134.5	39926.3	31081.8	34898.5	31438.6	33790.7
OTUB1_HUMAN	Ubiquitin thioesterase OTUB1 OS=Homo sapiens GN=OTUB1 PE=1 SV=2	7	6	1.0	0.933		859.1	882.3	668.4	11999.2	22954.0	14320.8	22107.3	14942.4	12217.7	14220.7	6664.5	7749.3	20701.0	5153.9	25780.9	5863.7	5753.4	6120.2
PCNA_HUMAN	Proliferating cell nuclear antigen OS=Homo sapiens GN=PCNA PE=1 SV=1	13	12	1.0	0.872		12855.7	12907.0	13179.7	15899.1	20630.4	38383.0	29624.5	28259.5	35594.5	19830.8	24331.2	18124.3	26388.6	31521.8	22544.9	20065.3	19475.0	19602.0
TBA1C_HUMAN	Tubulin alpha-1C chain OS=Homo sapiens GN=TUBA1C PE=1 SV=1	55	1	1.0	0.985		10.5	0.0	7.0	0.0	0.0	0.0	809.4	0.0	517.4	262.1	937.1	0.0	59.2	0.0	0.0	34.8	8.2	18.7
PRP19_HUMAN	Pre-mRNA-processing factor 19 OS=Homo sapiens GN=PRPF19 PE=1 SV=1	7	4	1.0	0.997		3720.8	4102.9	3964.3	27083.5	28401.3	17695.3	150267.0	167937.3	161579.2	141111.8	161368.9	144902.1	30855.5	39528.1	34888.9	3973.6	3550.2	3599.3
HS71L_HUMAN	Heat shock 70 kDa protein 1-like OS=Homo sapiens GN=HSPA1L PE=1 SV=2	61	3	1.0	0.991		554.0	390.3	588.2	12825.5	6925.6	14805.3	17955.5	18155.7	19662.2	10747.5	15097.7	11132.7	17944.7	15540.1	18047.2	1372.7	1119.2	1254.2
CTCFL_HUMAN	Transcriptional repressor CTCFL OS=Homo sapiens GN=CTCFL PE=1 SV=2	7	3	1.0	0.963		846.3	955.5	856.4	120070.3	121098.2	113885.5	126013.8	134246.6	149405.6	76494.2	78836.5	78339.2	145235.1	188057.2	165221.8	16576.4	16044.4	15744.5
HDAC8_HUMAN	Histone deacetylase 8 OS=Homo sapiens GN=HDAC8 PE=1 SV=2	207	180	1.0	0.876		5211537.1	5281044.7	5490803.0	11940447.7	12326642.8	11832059.2	14502516.8	16549191.6	16978024.1	17556600.2	17130364.0	16094406.6	10114861.9	11912184.2	11781740.2	6172088.6	6167883.9	6334716.6
DX39A_HUMAN	ATP-dependent RNA helicase DDX39A OS=Homo sapiens GN=DDX39A PE=1 SV=2	9	1	1.0	0.946		677.6	541.8	712.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	696.9	619.0	712.8
ATPA_HUMAN	ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 PE=1 SV=1	27	22	0.9	0.913		24810.7	24367.0	26998.3	160275.2	170369.7	221522.6	1138090.2	1212458.5	1068659.2	865245.7	974822.4	913179.5	414915.5	525925.5	449455.4	39013.1	40748.8	40004.2
ENOB_HUMAN	Beta-enolase OS=Homo sapiens GN=ENO3 PE=1 SV=5	13	3	0.9	0.839		295.9	323.5	283.1	6290.6	4099.9	6053.6	12634.4	9272.3	9264.0	2681.1	5535.5	5105.7	13094.3	10513.9	12784.9	1064.5	993.1	934.8
RS3A_HUMAN	40S ribosomal protein S3a OS=Homo sapiens GN=RPS3A PE=1 SV=2	21	20	0.9	0.765		13359.8	12715.5	12710.9	108480.7	145043.5	154032.0	129577.3	187062.3	159698.7	96231.5	93187.6	82879.9	147238.9	229676.1	138116.9	72936.0	68511.5	74162.7
ALDOB_HUMAN	Fructose-bisphosphate aldolase B OS=Homo sapiens GN=ALDOB PE=1 SV=2	3	1	0.9	0.909		0.0	98.5	96.8	0.0	0.0	0.0	0.0	1348.7	1249.5	0.0	0.0	0.0	1209.7	608.9	968.6	149.1	46.1	67.3
MDHC_HUMAN	Malate dehydrogenase, cytoplasmic OS=Homo sapiens GN=MDH1 PE=1 SV=4	10	8	0.9	0.788		4444.4	4030.0	4494.9	8554.6	7067.3	12140.8	19762.6	23884.0	23116.4	12654.0	17932.4	10569.9	33188.8	7751.8	21294.8	5150.4	4577.8	4752.7
RPN2_HUMAN	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2 OS=Homo sapiens GN=RPN2 PE=1 SV=3	4	3	0.9	0.820		924.8	1033.1	1052.6	63054.8	56470.7	45200.5	77509.1	91025.4	123988.0	40947.2	79400.0	72137.4	55922.3	161384.9	74658.4	6804.5	7581.6	7391.5
EFHD1_HUMAN	EF-hand domain-containing protein D1 OS=Homo sapiens GN=EFHD1 PE=1 SV=1	1	1	0.9	0.849		332.7	297.4	250.3	0.0	0.0	0.0	296.2	0.0	0.0	117.8	0.0	0.0	0.0	167.4	83.7	352.2	320.9	256.3
HNRPD_HUMAN	Heterogeneous nuclear ribonucleoprotein D0 OS=Homo sapiens GN=HNRNPD PE=1 SV=1	4	3	0.9	0.783		737.1	849.4	1333.6	22.8	18.2	39.8	1538.3	181.4	118.4	319.7	72.5	320.9	461.8	1695.9	423.6	883.2	577.0	726.1
RL10A_HUMAN	60S ribosomal protein L10a OS=Homo sapiens GN=RPL10A PE=1 SV=2	3	3	0.9	0.858		16.3	5.9	5.3	0.0	844.6	184.0	69.7	0.0	0.0	148.7	290.6	332.3	239.9	107.6	157.5	3.2	3.3	9.8
EIF3G_HUMAN	Eukaryotic translation initiation factor 3 subunit G OS=Homo sapiens GN=EIF3G PE=1 SV=2	2	2	0.9	0.706		533.9	366.0	493.7	6503.5	6373.9	5234.6	12181.2	14150.6	11464.5	11818.4	7396.2	12404.7	14666.7	6754.6	9633.2	813.1	1217.2	1213.0
MARE1_HUMAN	Microtubule-associated protein RP/EB family member 1 OS=Homo sapiens GN=MAPRE1 PE=1 SV=3	4	3	0.9	0.838		621.6	680.2	547.0	14079.3	0.0	1097.8	1079.5	369.2	1604.9	958.9	1212.7	2732.8	400.4	1993.0	181.7	5361.2	5187.8	5151.2
SYDC_HUMAN	Aspartate--tRNA ligase, cytoplasmic OS=Homo sapiens GN=DARS PE=1 SV=2	4	3	0.9	0.797		1037.0	1168.0	1251.6	6941.8	944.0	1250.9	32779.4	32626.6	33299.4	39824.8	39642.6	27114.9	10213.7	5619.5	3767.9	1090.8	993.8	798.7
NUDC_HUMAN	Nuclear migration protein nudC OS=Homo sapiens GN=NUDC PE=1 SV=1	8	6	0.8	0.707		4848.2	5304.6	4932.1	15082.3	15354.7	20157.6	22593.4	24069.0	29372.7	2901.6	6370.9	2524.2	36386.9	49678.3	48890.9	6798.7	7084.9	6746.3
U2AF2_HUMAN	Splicing factor U2AF 65 kDa subunit OS=Homo sapiens GN=U2AF2 PE=1 SV=4	1	1	0.8	0.899		0.0	3.5	13.3	0.0	0.0	0.0	3402.2	0.0	0.0	0.0	0.0	4110.4	0.0	0.0	0.0	0.0	0.0	0.0
RL19_HUMAN	60S ribosomal protein L19 OS=Homo sapiens GN=RPL19 PE=1 SV=1	3	3	0.8	0.567		761.9	906.1	908.8	734965.5	661079.5	651433.7	612256.9	567158.9	798744.8	364860.5	228958.3	305019.6	1161882.0	1456381.5	1358811.3	100337.4	115210.8	110590.7
RS2_HUMAN	40S ribosomal protein S2 OS=Homo sapiens GN=RPS2 PE=1 SV=2	8	8	0.8	0.524		1873.6	1825.0	1882.1	68153.2	56152.2	49309.9	75257.5	81290.3	108995.2	43666.5	36006.9	18967.9	129488.5	153973.0	149804.3	18133.8	18375.7	17891.9
RL13A_HUMAN	60S ribosomal protein L13a OS=Homo sapiens GN=RPL13A PE=1 SV=2	1	1	0.7	0.431		262.8	381.1	286.0	492.8	882.4	2041.2	1492.5	1965.7	1771.7	811.4	1464.0	942.2	2678.4	3377.7	2177.2	538.1	422.6	376.7
HPTR_HUMAN	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=2 SV=2	2	1	0.7	0.661		47.3	11.1	0.0	9537.7	3662.3	300.9	3670.4	35908.3	16198.4	27646.1	32478.7	23516.4	2823.8	2483.3	2801.1	502.8	541.6	550.7
CALR_HUMAN	Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1	12	12	0.7	0.503		5871.6	6719.7	5886.5	21346.1	18491.8	23649.0	156000.5	183740.4	210757.8	168816.3	225948.0	146204.2	94483.3	111852.8	94059.3	9866.2	10181.6	8813.3
GBB4_HUMAN	Guanine nucleotide-binding protein subunit beta-4 OS=Homo sapiens GN=GNB4 PE=1 SV=3	2	1	0.7	0.208		36954.2	37681.7	39491.8	89481.7	139438.2	167633.5	171318.2	195616.5	178173.2	266433.5	288155.8	263777.8	175525.5	218800.7	201327.2	39167.3	42925.8	44443.6
SNX1_HUMAN	Sorting nexin-1 OS=Homo sapiens GN=SNX1 PE=1 SV=3	2	1	0.7	0.694		37.1	36.9	56.2	52.3	0.0	0.0	466.3	3434.3	0.0	0.0	3353.0	1117.5	1122.9	78.7	91.7	94.9	84.1	38.4
SPEE_HUMAN	Spermidine synthase OS=Homo sapiens GN=SRM PE=1 SV=1	3	1	0.7	0.226		5414.7	5049.1	5363.9	7341.6	2104.2	3686.0	11896.0	9103.0	13195.5	18595.2	17580.7	22455.7	5650.7	7177.0	6455.8	5620.0	5017.5	4995.9
INO1_HUMAN	Inositol-3-phosphate synthase 1 OS=Homo sapiens GN=ISYNA1 PE=1 SV=1	1	1	0.7	0.603		187.2	205.8	142.2	0.0	0.0	0.0	4073.0	3696.5	6972.5	5261.8	9883.7	7683.3	0.0	0.0	13.2	0.0	5.4	0.0
RCC2_HUMAN	Protein RCC2 OS=Homo sapiens GN=RCC2 PE=1 SV=2	11	10	0.7	0.358		3836.0	4206.4	3869.6	577817.3	563470.6	407401.8	549182.6	551413.1	806183.5	357529.6	378281.7	293566.3	1119872.0	1262001.3	1480972.1	113796.6	127507.0	120578.9
AT1B3_HUMAN	Sodium/potassium-transporting ATPase subunit beta-3 OS=Homo sapiens GN=ATP1B3 PE=1 SV=1	1	1	0.7	0.516		160.7	116.6	42.7	0.0	0.0	0.0	0.0	114.2	409.1	441.3	149.3	461.3	39.2	0.0	81.7	22.1	47.7	47.9
RAB14_HUMAN	Ras-related protein Rab-14 OS=Homo sapiens GN=RAB14 PE=1 SV=4	4	4	0.6	0.350		4032.0	4417.3	3952.8	1910.2	497.2	0.0	6553.9	5194.5	9461.2	1203.4	342.1	277.9	16237.7	3544.6	18476.6	6004.3	5804.9	5435.9
PUR6_HUMAN	Multifunctional protein ADE2 OS=Homo sapiens GN=PAICS PE=1 SV=3	11	9	0.6	0.344		4319.7	4019.7	2840.8	11004.0	42457.7	9857.3	63461.2	56228.4	84393.2	31111.5	18547.5	27392.2	115608.5	113576.1	105737.3	11805.4	10526.0	10661.2
STRA8_HUMAN	Stimulated by retinoic acid gene 8 protein homolog OS=Homo sapiens GN=STRA8 PE=2 SV=1	1	1	0.6	0.343		2627.5	2230.8	2662.0	3341.2	937.0	0.0	48436.2	49097.4	49987.5	46469.4	53691.6	65486.8	21332.9	32279.7	27260.1	3234.4	3406.1	3280.4
SCMC1_HUMAN	Calcium-binding mitochondrial carrier protein SCaMC-1 OS=Homo sapiens GN=SLC25A24 PE=1 SV=2	1	1	0.6	0.302		703.3	746.1	615.9	0.0	0.0	0.0	1072.0	1328.4	1426.3	241.1	83.1	504.8	2730.4	1885.6	2440.6	587.2	522.7	607.8
LDH6A_HUMAN	L-lactate dehydrogenase A-like 6A OS=Homo sapiens GN=LDHAL6A PE=2 SV=1	5	3	0.6	0.035		34844.9	33213.9	35804.4	7079.8	3477.5	3792.2	22914.1	21080.4	22686.2	41955.7	23919.2	18277.7	28988.8	27921.2	25169.1	45645.4	45508.7	45946.4
UBXN1_HUMAN	UBX domain-containing protein 1 OS=Homo sapiens GN=UBXN1 PE=1 SV=2	1	1	0.6	0.445		438.7	402.8	461.7	0.0	3409.3	7132.1	1882.7	6273.7	10217.5	10863.9	25956.8	8041.9	570.7	1888.1	2991.7	759.7	495.9	512.3
ENPLL_HUMAN	Putative endoplasmin-like protein OS=Homo sapiens GN=HSP90B2P PE=5 SV=1	8	2	0.5	0.389		426.3	391.1	300.2	0.0	67.5	0.0	1219.0	4226.0	543.7	441.3	6007.2	2757.1	1837.5	110.0	1752.4	180.3	157.2	233.0
ACTY_HUMAN	Beta-centractin OS=Homo sapiens GN=ACTR1B PE=1 SV=1	3	1	0.5	0.089		5534.6	5836.4	6531.0	612.2	0.0	0.0	9034.3	7890.9	7982.9	14819.4	852.2	17171.5	12658.6	15549.8	6039.2	5193.4	4561.9	4912.7
POTEI_HUMAN	POTE ankyrin domain family member I OS=Homo sapiens GN=POTEI PE=3 SV=1	30	1	0.5	0.418		288.1	165.6	143.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	403.6	330.4	418.1
NSF1C_HUMAN	NSFL1 cofactor p47 OS=Homo sapiens GN=NSFL1C PE=1 SV=2	4	3	0.5	0.254		648.8	835.2	717.8	0.0	0.0	0.0	2006.2	664.7	1459.3	0.0	0.0	0.0	4233.7	3663.7	2852.2	628.8	823.0	692.0
CPSF5_HUMAN	Cleavage and polyadenylation specificity factor subunit 5 OS=Homo sapiens GN=NUDT21 PE=1 SV=1	1	1	0.5	0.416		0.9	0.0	2.3	0.0	0.0	0.0	0.0	133.1	0.0	0.0	0.0	0.0	94.7	19.3	106.4	6.8	23.1	37.3
TBB2B_HUMAN	Tubulin beta-2B chain OS=Homo sapiens GN=TUBB2B PE=1 SV=1	54	1	0.4	0.112		245.9	285.6	207.2	0.0	305.1	2451.0	3375.7	5045.0	0.0	5516.5	7132.9	5261.4	2891.1	3391.9	2566.9	214.3	386.5	371.9
FKBP3_HUMAN	Peptidyl-prolyl cis-trans isomerase FKBP3 OS=Homo sapiens GN=FKBP3 PE=1 SV=1	3	2	0.4	0.112		351.3	422.8	436.5	2875.9	354.7	3266.1	787.1	852.0	470.6	403.7	1767.1	4657.3	6196.1	3666.2	5193.9	458.5	471.9	472.6
EF1A2_HUMAN	Elongation factor 1-alpha 2 OS=Homo sapiens GN=EEF1A2 PE=1 SV=1	31	2	0.3	0.172		378.9	454.1	417.0	2977.0	14251.1	5838.5	17361.6	14527.3	17076.7	8837.4	5155.4	3306.2	57870.8	72438.3	64284.6	1219.2	1201.4	1044.3
CBR1_HUMAN	Carbonyl reductase [NADPH] 1 OS=Homo sapiens GN=CBR1 PE=1 SV=3	2	2	0.3	0.065		282.9	267.6	266.9	43.6	37.3	82.3	215.5	685.3	23.0	0.0	27.5	97.4	378.0	519.3	671.6	1427.9	1414.9	1224.2
H90B2_HUMAN	Putative heat shock protein HSP 90-beta 2 OS=Homo sapiens GN=HSP90AB2P PE=1 SV=2	30	4	0.3	0.289		179.1	172.1	162.0	3083.3	2871.2	5580.0	5261.7	5863.1	3799.0	1273.6	2946.8	1148.3	10366.2	56107.6	6780.4	3544.2	944.9	4390.1
EFTU_HUMAN	Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2	10	6	0.3	0.175		1560.6	1310.4	1751.0	1875.5	4237.4	16657.1	15618.3	4044.1	5638.2	4041.6	4115.1	2682.9	36528.1	85644.7	50996.0	2025.5	1870.2	1645.6
PCBP2_HUMAN	Poly(rC)-binding protein 2 OS=Homo sapiens GN=PCBP2 PE=1 SV=1	9	5	0.3	0.082		4485.9	4840.7	4899.5	9663.2	11336.9	12996.2	14197.2	10787.6	11039.5	11850.1	13457.6	12482.9	67601.1	106630.8	78825.4	7440.0	7964.5	8200.1
AL9A1_HUMAN	4-trimethylaminobutyraldehyde dehydrogenase OS=Homo sapiens GN=ALDH9A1 PE=1 SV=3	2	1	0.2	0.102		3.8	0.0	0.0	0.0	0.0	262.6	204.6	1588.1	663.4	127.8	235.7	622.8	2022.0	4031.2	3390.2	217.0	322.0	412.1
RL14_HUMAN	60S ribosomal protein L14 OS=Homo sapiens GN=RPL14 PE=1 SV=4	3	3	0.2	0.053		938.5	886.8	949.3	0.0	0.0	0.0	609.7	434.3	854.0	333.0	223.0	611.8	6519.8	6702.0	5772.0	1162.6	1309.2	1151.1
MAOM_HUMAN	NAD-dependent malic enzyme, mitochondrial OS=Homo sapiens GN=ME2 PE=1 SV=1	1	1	0.2	0.220		18.3	9.1	17.1	0.0	0.0	247.2	445.2	0.0	339.2	1839.6	3074.2	442.6	0.0	111.0	0.0	0.0	8.6	25.6
RL15_HUMAN	60S ribosomal protein L15 OS=Homo sapiens GN=RPL15 PE=1 SV=2	1	1	0.2	0.078		0.0	8.0	6.3	0.0	0.0	0.0	1019.7	1025.9	1638.1	681.0	84.5	0.0	5564.1	7293.6	5597.0	777.5	639.4	752.9
1433S_HUMAN	14-3-3 protein sigma OS=Homo sapiens GN=SFN PE=1 SV=1	4	2	0.0	0.059		0.0	0.0	1.1	0.0	0.0	0.0	119.5	448.7	128.1	0.0	196.9	0.0	21935.9	21764.0	19422.3	1845.9	1282.1	1482.0
PSA5_HUMAN	Proteasome subunit alpha type-5 OS=Homo sapiens GN=PSMA5 PE=1 SV=3	2	1	0.0	0.139		1.9	2.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	224.6	0.0	251.0	0.0	0.0	9.6	6.8	20.6
TEG5_HHV1F	Tegument protein UL47 OS=Human herpesvirus 1 (strain F) GN=UL47 PE=3 SV=2	4	2	0.0	0.061		131.3	1007.2	36.7	0.0	0.0	0.0	216.4	0.0	0.0	0.0	0.0	18.5	212846.8	318398.4	268802.9	16235.8	1172.0	45188.8
PLSL_HUMAN	Plastin-2 OS=Homo sapiens GN=LCP1 PE=1 SV=6	10	1	0.0	0.084		0.0	1.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	514.1	594.2	28.9	41.6	207.3	12.3	0.0	0.0