Table S1. High-throughput next-generation RNA sequencing of non-malignant HCEC-1CT and malignant HCT116 and DLD-1 cell lines originating from colon tissue. In the blue-colored table are presented FPKM values for the whole expression of the *SMAD4* gene, while in the green-colored table are presented FPKM values for each SMAD4 transcript.

gene_id	gene_name	gene_locus
ENSG00000141646	SMAD4	18:51028394-51085045

transcript_id	transcript_name	gene_id	gene_name	transcript_locus
ENST00000589076	SMAD4-207	ENSG00000141646	SMAD4	18:51030100-51048708
ENST00000590061	SMAD4-210	ENSG00000141646	SMAD4	18:51030213-51048857
ENST00000592911	SMAD4-215	ENSG00000141646	SMAD4	18:51030597-51058154
ENST00000588745	SMAD4-205	ENSG00000141646	SMAD4	18:51047047-51078467
ENST00000592186	SMAD4-214	ENSG00000141646	SMAD4	18:51047047-51078467
ENST00000342988	SMAD4-201	ENSG00000141646	SMAD4	18:51030213-51085039
ENST00000398417	SMAD4-202	ENSG00000141646	SMAD4	18:51029614-51085045
ENST00000591126	SMAD4-212	ENSG00000141646	SMAD4	18:51052326-51079777
ENST00000611848	SMAD4-217	ENSG00000141646	SMAD4	18:51058353-51079143
ENST00000591914	SMAD4-213	ENSG00000141646	SMAD4	18:51030213-51049324
ENST00000589941	SMAD4-209	ENSG00000141646	SMAD4	18:51044959-51048798
ENST00000589706	SMAD4-208	ENSG00000141646	SMAD4	18:51047179-51049565
ENST00000588860	SMAD4-206	ENSG00000141646	SMAD4	18:51028394-51048816
ENST00000590499	SMAD4-211	ENSG00000141646	SMAD4	18:51066822-51077291
ENST00000586253	SMAD4-204	ENSG00000141646	SMAD4	18:51077251-51078870
ENST00000593223	SMAD4-216	ENSG00000141646	SMAD4	18:51067113-51077130
ENST00000585448	SMAD4-203	ENSG00000141646	SMAD4	18:51051093-51054916

HCT116 (FPKM)	DLD-1 (FPKM)	HCEC-1CT (FPKM)
30.884188	43.799004	30.83934

HCT116 (FPKM)	DLD-1 (FPKM)	HCEC-1CT (FPKM)
0	0	0
0	0	0
0	0	0
0	0	0
0	0	0
3.168145	1.815537	2.689202
0.013276	0.001923	0.038767
4.459052	9.533424	6.472704
1.792871	2.256931	1.294333
11.492303	11.510384	1.571642
0.036732	0.158341	4.97713
0.330983	3.068607	2.06937
0	0	0
0.288089	1.159342	0.478574
0.275016	0.487777	0.344155
0.515416	1.476431	0.664413
0	0	0

Table S2. Diferentially bound transcriptional regulators to SMAD4 promoters C and D predicted by bioinformatic tools Tfsitescan, AliBaba2.1, PROMO, TFBIND and CiiiDER.

Promoter C	Promoter D
ADF1	OCT1
AHR	AFP1
AP2	ALX1
AP3	ALX4
ARNT::HIF1A	ANTP
CMYC	AP1
CPE_BINDING_PRO	AP1FJ
CTCFL	ARID3B
E1AF	ARID5A
E2F1	ARX
E2F2	ASCL1
E2F4	ATF3
E2FDRTF	ATOH1
E4F1	ATOH7
EBF1	BATF
EBF3	BATF::JUN
EGR1	BATF3
EGR3	BHLHA15
EGR4	BRACH
EHF	BRN2
ELF1	BSX
ELK1	C/EBP
ELK3	C/EBPBETA
ELK4	C/EBPDELTA
ETF	CDPCR1
ETS1	CDPCR3HD
ETS2	CDX4
ETV4	CEBPALPHA
FEV	CF1
FOXD1	CIZ8
FOXE1	CJUN
FOXK1	COUP
FOXK2	CREB
FOXO6	CUX1
FOXP2	CUX2
FOXQ1	DBP
GABPA	DDIT3::CEBPA
GCF	DL
GCM1	DLX1
GLI3	DLX3

HAND1::TCF3	DLX4
HEN1	DMRTA2
HES1	DPRX
IRF2	DRGX
ISL1	E4BP4
ISL2	ELF5
KLF1	EN1
KLF4	EOMES
KLF5	FOSB::JUNB
KLF9	FOSL1::JUNB
KROX20	FOSL1::JUND
LVC	FOXA2
MAZ	FOXA3
MEF2C	FOXB1
MEP1	FOXC1
MSANTD3	FOXC2
MTBF	FOXH1
NET_SAP2	FOXI1
NF1B1	FOXJ2
NFIC	FOXJ3
NFKAPPAB1	FOXN3
NFMUE1	FOXO1
NKX23	FOXP1
NKX25	FTZ
NKX3.2	GATA1
NKX31	GATA2
NKX32	GATA3
NR1I3	GBX1
NR4A2	GBX2
NRF1	GCN4
OCT1A	GFI1
OSR1	GFI1B
OSR2	GLO
OVOL1	GMEB1
OVOL2	GMEB2
P40X	GRE
PAX4PD	GSX1
PAX5	H3ABP
PEA2_2	HESX1
PLAGL1	HFH2
POU3F2	HIC1
POU3F3	HLF
POU5F1B	HLTF
PPARALPHA:RXRAL	HMBOX1
R2	HMX2
RAP1	HNF1A

RARALPHA1	HNF1B
RARBETA:RXRALPH	HNF1C
REB1	HNF3B
RELA	HNF4A
RELB	HNF4G
REPRESSOR_OFA	HOXA10
RORB	HOXA11
RREB1	HOXA2
RUNX3	HOXA6
SIF	HOXA7
SOX12	HOXB5
SP1	HOXB8
SP3	HOXB9
SP8	HOXC10
SPDEF	HOXC11
STAT5A	HOXC12
TCF12	HOXC13
TCF3	HOXC9
TCF7L1	HOXD10
TCFL5	HOXD11
TFAP2A	HOXD12
TFAP2B	HOXD3
TFAP2C	HOXD9
VEZF1	HSEBINDING_PRO
WT1	ICSBP
YY2	IK3
ZBTB14	ISGF3
ZBTB33	ISX
ZBTB6	JUN::JUNB
ZBTB7A	LBX2
ZBTB7C	LHX1
ZFP57	LHX3
ZFX	LHX6
ZIC1	LHX9
ZIC3	LIN54
ZIC4	LMX1A
ZNF219	LMX1B
ZNF423	LVB
ZNF460	LYF1
	MAFB
	MAX
	MEF2A
	MEIS2
	MEOX2
	MGA
	MIXA

MIXL1	
MNX1	
MSGN1	
MSX2	
MSX3	
MUEBPC2	
MYBL1	
NFATC1	
NFATC4	
NFBA1	
NFE2L2	
NFIX	
NKX25	
NKX28	
NKX3_1R52C	
NKX61	
NKX63	
NOBOX	
NOTO	
NR2C2	
NR2E1	
NR4A1	
NR5A1	
NRL	
NRSF	
OCT2.1	
OTX1	
PAX6	
PDX1	
PIT1A	
PITX1	
PITX2	
PITX3	
POU2F2	
POU6F1	
POU6F2	
RAX2	
RBPJKAPPA	
RHOX11	
RORA2	
RUSH	
SCRT1	
SCRT2	
SEF1	
SHOX2	
SIX1	

SMAD2::SMAD3
SNAI1
SNAI2
SOX13
SOX14
SOX15
SOX17
SOX2
SOX3
SOX8
SOX9
STAT1::STAT2
STAT2
TAL1ALPHAE47
TAL1BETAE47
TAL1BETAITF2
TBP
TBR1
TBX1
TBX2
TBX3
TBX4
TCF7L2
TEC1
TFAP4
TWIST2
UNCX
VBP
VMAF
VMW65
XBP1
XFD1
ZBTB12
ZEN1
ZKSCAN1
ZNF217
ZNF354C

Table S3. Proteomics analysis of transcriptional regulators interacting with promoter regions

Protein.Group	Protein.Ids	Protein.Names	Genes	HCEC1CT_(	HCEC1CT_	HCEC1CT_N	SW480_C	SW480_D	SW480_NC	SW620_C	SW620_D	SW620_NC
A0A024R1R8;Q9Y2S6	Q9Y2S6;A0A024R1R8	TMA7B_HUMAN;TMA7_HUMAN	TMA7;TMA7B				3912.09	548.562			3760.8	
A0A075B6H7;A0A0C4DH55;P0162	A0A075B6H7;A0A0C4DH90;A0A0C4DH55;P01624	KV315_HUMAN;KV37_HUMAN;KVD07_HUM	IGKV3-15;IGKV3-7;IGKV3D-7									
A0A0B4J2D5;P0DPI2	P0DPI2;A0A0B4J2D5	GAL3A_HUMAN;GAL3B_HUMAN	GATD3;GATD3B						3307.97			
A0A0C4DH36	A0A0C4DH36	HV338_HUMAN	IGHV3-38	3366.45			2239.52	1139.11	1193.9			
A0AV96	A0AV96	RBM47_HUMAN	RBM47					4957.72		878.12	10243.4	5718.77
A0AVT1	A0AVT1	UBA6_HUMAN	UBA6					3300.5	3384.65	3090.23		
A0FGR8	A0FGR8	ESYT2_HUMAN	ESYT2	3744.35	3037.41	5838.97	3463.32	6437.14	4607.32	16105.9	5262.65	11386.9
A0MZ66	A0MZ66	SHOT1_HUMAN	SHTN1						872.127	1117.13	612.393	1082.73
A1L0T0	A1L0T0	HACL2_HUMAN	ILVBL		3183.67		1839.74	3269.61	4218.14	4285.24	3258.41	3600.14
A1L390	A1L390	PKHG3_HUMAN	PLEKHG3					3813.71	2069.52	11804.2		8161.37
A2RRP1	A2RRP1	NBAS_HUMAN	NBAS		997.991		972.653	1788.9	936.137	1439.25	1061.94	716.029
A3KMH1	A3KMH1	VWA8_HUMAN	VWA8	3401.23	2447.23	3620.95	3835.52	4495.24	3793.46	4114.52	3906.2	3788.36
A5YKK6	A5YKK6	CNOT1_HUMAN	CNOT1	5826.81	4832.17	6319.4	7985.91	5579.34	5700.35	7827.39	8087.22	7570.27
A6NHL2	A6NHL2	TBAL3_HUMAN	TUBAL3	4017.02	2351.02	2353.52	3331.23	1649.45		4070.88	3179.3	4912.77
A6NHR9	A6NHR9	SMHD1_HUMAN	SMCHD1	1915.97	1335.46	1425.82	1638.7	2349.65	2384.61	4606.64	1153.86	4287.43
A6NJ78	A6NJ78	MET15_HUMAN	METTL15			1832.28	2524.56	2547.06	1772.65	2271.28	2134.43	2800.83
A7E2V4	A7E2V4	ZSWM8_HUMAN	ZSWIM8		4178.97		1783.41	2644.49	1742.23	1341.63	1847.73	2354.47
A7KAX9	A7KAX9	RHG32_HUMAN	ARHGAP32							2904.66		3242.06
A8K2U0	A8K2U0	A2ML1_HUMAN	A2ML1			5000.6	3075.81		2161.16	1663.22	2468.79	2472.46
A8MWD9;P62308	A8MWD9;P62308	RUXGL_HUMAN;RUXG_HUMAN	SNRPG;SNRPGP15				4631.27	3522.3	1748.43	2905.09	8285.89	3090.33
B5ME19;Q99613	Q99613;B5ME19	EIF3C_HUMAN;EIF3CL_HUMAN	EIF3C;EIF3CL	11273.8	9152.91	10113.7	31143.6	18356	23284.6	22413.8	13852.8	17343.3
B9A064;P0CG04	P0DOY3;P0CF74;P0DOY2;B9A064;P0CG04;A0M8Q6	IGLC1_HUMAN;IGLL5_HUMAN	IGLC1;IGLL5	7755.68	4014.98	16783.3	4606.89	2862.19	1984.3	3318.67	5421.99	2935.48
C9JN71	C9JN71	ZN878_HUMAN	ZNF878	15906.3	12614.1	27135.8	6140.17	82885.2	41629.3	46807.2	15646.3	51768.2
C9JRZ8	C9JRZ8	AK1BF_HUMAN	AKR1B15									
E9PAV3	E9PAV3	NACAM_HUMAN	NACA	5816.59		3414.06	6113.21	6427.77	12197.5	5970.33	8330.1	4662.23
E9PRG8	E9PRG8	CK098_HUMAN	C11orf98				1067.53	5282.31	2824.05	1157.12	1838.14	1328.83
L0R819	L0R819	ASURF_HUMAN	ASDURF						1328.38	2358.44		2247.89
O00116	O00116	ADAS_HUMAN	AGPS				6667.37	7746.93	4649.5	4848.19	4633.53	4426.85
O00139	O00139	KIF2A_HUMAN	KIF2A	3113.38	4465.88	3215.25	2904.16	4814	4358.48	2975.27	1394.49	3929.22
O00148	Q13838;O00148	DX39A_HUMAN	DDX39A	37065	17437.6	33474.6	54332	34888.2	34843	51756.8	63405	45559.4
O00151	O00151	PDLI1_HUMAN	PDLIM1	17078.5	3470.77	15021.5	6955.95	8242.18	17046.8	1860.72	61434.9	2858.57
O00154	O00154	BACH_HUMAN	ACOT7	4145.93	3641.01	5154.91	4512.3	4298.95	6288.06	4884.39	2514.21	3335.21
O00159	O00159	MYO1C_HUMAN	MYO1C	42363.8	41480.4	41038	24522	33821.6	23526.5	187277	69217.3	138733
O00161	O00161	SNP23_HUMAN	SNAP23	3702.23	5299.14	4230.72	21155.5	52926.5	58516.1	86889.4	10689.6	80737.5
O00165	O00165	HAX1_HUMAN	HAX1				3982.59	769.787	1807.84	2274.68	5965	8728.16
O00170	O00170	AIP_HUMAN	AIP	4399.82	2006.63	4240.11	7473.25	3036.2	2249.69	6139.69	2880.92	5784.45
O00178	O00178	GTPB1_HUMAN	GTPBP1		1187.09	1455.22	1772.99	2287.98	2179.89	1103.06		1339.52
O00182	O00182	LEG9_HUMAN	LGALS9							2408.85		2347.49
O00186	O00186	STXB3_HUMAN	STXBP3	1208.11	1107.4		1582.13	3438.82	2388.95	1399.19	898.882	1737.2
O00203	Q13367;O00203	AP3B1_HUMAN	AP3B1	5435.87	6302.26	4133.51	7246.42	5756.16	4298.9	3780.78	5289.88	2554.56
O00213	O00213	APBB1_HUMAN	APBB1					2233.29	1968.74			
O00214	O00214	LEG8_HUMAN	LGALS8				5525.12	6848.34	6542.16	26201.8	8511.46	24288.6
O00231	O00231	PSD11_HUMAN	PSMD11	15388.9	16196.5	16469	17749.4	19223.9	20032.3	21047.8	12093.4	15987.5
O00232	O00232	PSD12_HUMAN	PSMD12	11015.5	9855.71	12409.8	18232.5	17585.3	18903.9	14007.7	8804.9	15261.1
O00233	O00233	PSMD9_HUMAN	PSMD9	9906.54	8331.87	10665.6	9532.76	3893.9	7383.89	6607.62	9526.59	7499.04
O00267	O00267	SPT5H_HUMAN	SUPT5H				1502.93	1182	1917.05	1872.25		991.885
O00273	O00273	DFFA_HUMAN	DFFA				682.896		3882.89	2523.25	753.402	1636.42
O00291	O00291	HIP1_HUMAN	HIP1	13662		17792.1				23983.2		21707.8
O00299	O00299	CLIC1_HUMAN	CLIC1	14550.1	12458.4	17746.5	17088.4	6969.5	12223.5	4454.81	8432.73	5182.42
O00303	O00303	EIF3F_HUMAN	EIF3F	4272.8		4738.85	7454.24	15869.6	4993.07	7994.83	4764.73	6727.04
O00330	O00330	ODPX_HUMAN	PDHX	5677.21	5770.59	11537	42520.1	2196.91	12526.9	2447.96	3144.54	5416.4
O00401	O00401	WASL_HUMAN	WASL			2726.29		1515.66	2548.99			
O00410	O00410	IPO5_HUMAN	IPO5	8699.52	7719.02	8934.75	13480.8	13574	11884.2	16896	15580.6	13953.5
O00411	O00411	RPOM_HUMAN	POLRMT							1999.03	3377.75	2424.91
O00425	O00425	IF2B3_HUMAN	IGF2BP3	16644.8	8259.91	6952.73	21363.3	53328.1	15066.7	7894.5	10635.2	11558.6
O00429	O00429	DNM1L_HUMAN	DNM1L	2968.82	2899.9	3553.2	12562	6211.3	9364.95	7503.02	11527.7	6645.67
O00442	O00442	RTCA_HUMAN	RTCA	3123.49		1998.48	3744.63	7656.63	5334.47	3835.18	3041.31	4743.81
O00443	O00443	P3C2A_HUMAN	PIK3C2A				1482.23	2889.57	2997.3	1888.1	3192.6	2210.53
O00458	O00458	IFRD1_HUMAN	IFRD1					3608.14	1113.98	582.02		808.86
O00461	O00461	GOLI4_HUMAN	GOLIM4				428.6	1318.71	1285.23	2429.94		1970.37
O00463	O00463	TRAF5_HUMAN	TRAF5							1573.31		1964.81
O00469	O00469	PLOD2_HUMAN	PLOD2	2115.63		3384.52	3939.32	2163.27	2048.9	1445.61		
O00471	O00471	EXOC5_HUMAN	EXOC5		1475.61		1137.73	1175.22	999.96	880.86	1227.62	1116.35
O00483	O00483	NDUA4_HUMAN	NDUFA4				7082.09				12563.3	
O00487	O00487	PSDE_HUMAN	PSMD14		5118.68	7171.9	9319.3	9604.98	8372.1	8513.48		8089.47
O00499	O00499	BIN1_HUMAN	BIN1				1313.5		1988.81			
O00505	O00629;O00505	IMA4_HUMAN	KPNA3	3475.59		3723.33	2399.07	3896.12	4162.22	4306.7	1907.67	4590.64
O00515	O00515	LAD1_HUMAN	LAD1							11593	6978.18	9170.3
O00541	O00541	PESC_HUMAN	PES 1.00	3766.79	2731.72	3976.28	2245.37	6641.89	4695.95	4288.35	2923.63	4322.85
O00560	O00560	SDCB1_HUMAN	SDCBP	6617.26	2188.38	5711.24	3664.19	2166.48	2304.33	1453.86	1872.1	1021.85
O00567	O00567	NOP56_HUMAN	NOP56				2789.78	2924.67	1565.92	2383.12	3897	3017.6
O00571	O15523;O00571	DDX3X_HUMAN	DDX3X	20042	19014.1	18357.2	17230.1	25533.1	16556.7	14503.1	9577.19	17348.2
O00629	O00629	IMA3_HUMAN	KPNA4				709.569	1308.13	1667.85	930.608		1102.13
O00712	O00712	NFIB_HUMAN	NFIB	14335.9								
O00743	O00743	PPP6_HUMAN	PPP6C			4532.06	4342.74	5484.19	9067.41	5485.86	4057.95	4567.4
O00763	O00763	ACACB_HUMAN	ACACB	86065.6	71054.2	58448.6	9860.33	32531.4	8561.62	35553.5	83987.9	38319.6
O00764	O00764	PDXK_HUMAN	PDXK						2433.75	1746.28		1749.85
O14493;O15551;O95484;P56747	P56747;O14493;O15551;O95484	CLD3_HUMAN;CLD4_HUMAN;CLD6_HUMA	CLDN3;CLDN4;CLDN6;CLDN9					13975.9	5445.98	20359.1	7568.68	15704.3
O14530	O14530	TXND9_HUMAN	TXNDC9				480.327	490.132	924.613	463.413		731.182
O14531	O14531	DPYL4_HUMAN	DPYSL4								4868.79	
O14556	O14556	G3PT_HUMAN	GAPDHS				2025.96	8059.46				
O14578	O14578	CTRO_HUMAN	CIT	46204	35773	28994.4	28717.8	10127.9	10018.5	16652.3	57241.4	12908.2
O14579	O14579	COPE_HUMAN	COPE	7656.65	8682.54	7353.34	6865.09	8166.43	8064.75	10772.8	9088.09	7934.37
O14617	O14617	AP3D1_HUMAN	AP3D1	4388.25	4248.9		1900.99	6626.9	3154.38	1647.24	1963.33	2705.6
O14639	O14639	ABLM1_HUMAN	ABLIM1					4113.88	1564.03	1535.58		
O14641	O14641	DVL2_HUMAN	DVL2					1180.34		1270.92	895.021	1456.07
O14646	O14647;O14646	CHD1_HUMAN	CHD1				1589.26	5561.17	3699.31	2756.7	1423.24	2564.8
O14653	O14653	GOSR2_HUMAN	GOSR2			2832.07	2752.88	2869.88	3046.59		1362.29	1887.8
O14656	O14656	TOR1A_HUMAN	TOR1A				2110.43	2832.59	2958.18	2326.9	2141.23	1462.86
O14672	O14672	ADA10_HUMAN	ADAM10	5271.8	3312.31	3969.82	4068.95	3083.6	5481.65	4683.1	2224.71	4460.89
O14730	O14730	RIOK3_HUMAN	RIOK3	3032.76		3412.11	5390.04	1395.94	1627.35	600.736	3375.6	4180.88
O14733	O14733	MP2K7_HUMAN	MAP2K7				24317.3	1980.21	1418.78	2042.21	29410.9	2437.59
O14734	O14734	ACOT8_HUMAN	ACOT8		6323.4	6609.79	3198.09	2311.69	1153.47	4923.08	3270.61	4787.81
O14735	O14735	CDIPT_HUMAN	CDIPT	4863.57	3481.37	7059.15	2670.32			2325.28	2627.83	2346.26
O14737	O14737	PDCD5_HUMAN	PDCD5	4392.69	1283.65	3194.75	4894.63	2717.33	8158.67	6315.77	5964.33	5215.84
O14744	O14744	ANM5_HUMAN	PRMT5	4962.45	3349.47	4439.66	5880.71	8683	9110.91	6293.13	4381.72	8652.63
O14745	O14745	NHRF1_HUMAN	NHERF1			2495.22	1688.6			1095.69	1985.12	1149.71
O14757	O14757	CHK1_HUMAN	CHEK1					2269.95	1903.42	3220.42	1701.2	2932.08
O14773	O14773	TPP1_HUMAN	TPP1	13673.2	10126.1	13954.2	9327.64	7845.42	5599.73	9239.7	15675.1	10211.2
O14776	O14776	TCRG1_HUMAN	TCERG1				783.182	2841.58	1659.49	1735.51		1532.14
O14802	O14802	RPC1_HUMAN	POLR3A	11366.7	14767.1	4496.38	2841.61	3869.42	1812.85	2957.99	2156	3019.69
O14818	O14818	PSA7_HUMAN	PSMA7	5149.09	4064.84	5713.71	4539.52	3280.26	6661.39	6187.82	4523.36	7129.77
O14828	O14828	SCAM3_HUMAN	SCAMP3	5408.52		5959.39	6673.64	4858.89	4317.78	3332.9	3668.3	2324.75
O14867	O14867	BACH1_HUMAN	BACH1	2732.88			2800.88	5189.56	3338.27			
O14874	O14874	BCKD_HUMAN	BCKDK				6703.83				5457.57	4776.84
O14879	O14879	IFIT3_HUMAN	IFIT3	3953.55	13220.6	2605.86	4161.43	4749.5	5321			
O14880	O14880	MGST3_HUMAN	MGST3	2252.62	3193.08			1850.44		1933.98	921.773	986.956
O14907	O14907	TX1B3_HUMAN	TAX1BP3						505.253	1076.48	1790.25	
O14908	O14908	GIPC1_HUMAN	GIPC1						2849.86			2068.63
O14929	O14929	HAT1_HUMAN	HAT1					2453.72	4239.52	3536.06		3878.32
O14939	O14939	PLD2_HUMAN	PLD2		2068.12	2434.97	1110.97	2782.84	2732.55	4656.63	1758.57	4425.35
O14950;P19105	P24844;P19105;O14950	ML12A_HUMAN;ML12B_HUMAN	MYL12A;MYL12B	41586.6	24828.5	31551.4	17483.8	59453.3	44833.7	60429.1	125830	34444.3
O14964	O14964	HGS_HUMAN	HGS	2907.62		3537.27	4866.64			1819.08	2738.36	
O14965	O14965	AURKA_HUMAN	AURKA				3827.24	4195.65	3364.94	3305.87	2363.86	3446.54
O14966;P57729;Q13637	O14966;P57729;Q13637	RAB32_HUMAN;RAB38_HUMAN;RAB7L_HU	RAB29;RAB32;RAB38			5768.82				3408.1	1708.51	2471.61
O14974	O14974	MYPT1_HUMAN	PPP1R12A	20642.1	18826.5	17394.8	53310.7	101261	65828	8171.44	9278.54	8210.18
O14976	O14976	GAK_HUMAN	GAK				1985.25	4260.46	4315.41	5586.33	3243.34	6607.34
O14979	O14979	HNRDL_HUMAN	HNRNPDL	3526.15			3359.51	16358.7	6802.63	3360.05	982.872	4338.2
O14980	O14980	XPO1_HUMAN	XPO1	15622.3	12410.9	12040.6	19177.3	13389.4	16724.8	25357.8	24647.5	24644.1
O14981	O14981	BTAF1_HUMAN	BTAF1	4363.25	4781.44		5318.13	6762.3	5181.87	5881.61	6146.78	5474.16
O15014	O15014	ZN609_HUMAN	ZNF609				1295.07	2978.85	2315.01	1657.86	931.814	1617.19
O15020	O15020	SPTN2_HUMAN	SPTBN2		582.102			3409.1	2272.09	1969.86	1412.06	1685.69
O15027	O15027	SC16A_HUMAN	SEC16A	4855.08	2501.73	2650.6	3732.41	4969.36	3663.72	4228.3	4824.14	5108.34
O15031	O15031	PLXB2_HUMAN	PLXNB2	4715.85	4665.76		4253.52	4766.83	4229.54	6446.51	5615.57	5525.45
O15042	O15042	SR140_HUMAN	U2SURP	2610.59	3520.24		3699.63	11358.6	4189.32	7810.86	4183.59	7217.95
O15056	O15056	SYNJ2_HUMAN	SYNJ2	17701	13547.4	10728.1	12920.5	8235.77	2959.69	4629.98	12042.2	6325.57
O15061	O15061	SYNEM_HUMAN	SYNM	13261.5	13819.8	10332.5		6972.62	5324.76			
O15067	O15067	PUR4_HUMAN	PFAS			4147.99	6882.59	2990.28	4458.72	8922.15	5485.73	8346.87
O15075	O15075	DCLK1_HUMAN	DCLK1									
O15084	O15084	ANR28_HUMAN	ANKRD28				1792.85	4343.38	3315.23	2383.16	1125.98	2595.6
O15091	O15091	MRPP3_HUMAN	PRORP		3777.78	3788.82	3589.86	5129.01	3403.55	9842.26	9435.54	9818.21
O15111	O15111	IKKA_HUMAN	CHUK	2263.13	2320.82	1731.4	4785.82	5951.48	5857.1	2815.59	2849.57	1929.29
O15117	O15117	FYB1_HUMAN	FYB1					9234.62	884.342			
O15118	O15118	NPC1_HUMAN	NPC1		2948.68			3674.77	3200.66	2800.02	2141.56	5171.42
O15119	O15119	TBX3_HUMAN	TBX3									13055.5
O15127	O15127	SCAM2_HUMAN	SCAMP2	4751.54		3207.47	6820.42	4411.52	5388.34	5095.43	4513.47	4562.47
O15143	O15143	ARC1B_HUMAN	ARPC1B	4559.24	2967.31	6445.17	3305.2	6562.7	11974.4	4764.64	1068.38	5227.15
O15144	O15144	ARPC2_HUMAN	ARPC2	8016.76	10839	13498.9	9497.99	16348.6	33980.9	12623.2	4441.08	14747.2
O15145	O15145	ARPC3_HUMAN	ARPC3		7739.95	11850.5	9264.89	19792	29513.1	11414.2	7028.57	8110.13
O15160	O15160	RPAC1_HUMAN	POLR1C	9254.19	14628.9		2827.93	3372.46	4934.42	5533.96	1595.58	3467.37
O15162	O15162	PLS1_HUMAN	PLSCR1					3382.25	3233.16	2574.73		
O15194	O15194	CTDSL_HUMAN	CTDSPL							2714.67		1442.02
O15212	O15212	PFD6_HUMAN	PFDN6	7746.25	8909.5	6708.77	4701.19	9459.19	7582.66	6629.88	4556.04	6562.19
O15228	O15228	GNPAT_HUMAN	GNPAT				4175.14		3944.62	3144.2	2943.1	4104.39
O15231	O15231	ZN185_HUMAN	ZNF185				340.221		1762.07			
O15234	O15234	CASC3_HUMAN	CASC3					3974.4	1975.34			
O15235	O15235	RT12_HUMAN	MRPS12	2102.42	3024.92	2557.75	2079		1894.12	1950.03	3221.11	1607.72
O15258	O15258	RER1_HUMAN	RER1	1406.45		3378.51	3727.87		1853.13	1816.49	2147.83	2564.23
O15269	O15269	SPTC1_HUMAN	SPTLC1	6938.78	9744.6	8452.25	7814.84	6325.09	5366.9	8531.21	10314.6	9612.88
O15270	O15270	SPTC2_HUMAN	SPTLC2				1535.8	1615.79	1121.79	1482.34	1227.4	1835.79
O15347	O15347	HMGB3_HUMAN	HMGB3	4275.22	3301.59		3146.1	4829.18	3910.73	3456.45	5902.05	3949.43
O15355	O15355	PPM1G_HUMAN	PPM1G				11728.4	10730.2	18571.3	13326.3	13039.7	13267.2
O15360	O15360	FANCA_HUMAN	FANCA					5017.3	2064.88	3956.7	4513.6	4032.64
O15371	O15371	EIF3D_HUMAN	EIF3D	3243.47	3125.91	3661.48	7256.18	6982.51	6570.21	4817.2	3717.84	5674.83
O15372	O15372	EIF3H_HUMAN	EIF3H	2140.52	1394.87	1955.23	4767.32	7294.93	8157.64	8860.91	3101.84	8812.39
O15382	O15382	BCAT2_HUMAN	BCAT2					1225.93	3040.05	1658.43		1593.15
O15397	O15397	IPO8_HUMAN	IPO8	5491.6	3155.94	4494.46	9288.02	2519	3392.59	7843.72	11979.9	10653.3
O15427	O15427	MOT4_HUMAN	SLC16A3	5011.91	4582.26	3886.12	2820.23	5579.28	2211.28	4937.47	5454.36	5320.05
O15439	O15439	MRP4_HUMAN	ABCC4							3471.62	1308.79	2449.29
O15460	O15460	P4HA2_HUMAN	P4HA2				4550.46	9549.09	4336.04	3669.43	3547.85	4961.16
O15479	O15479	MAGB2_HUMAN	MAGEB2				15630.2	37136	19562.3	1804.21	3168.11	2398.32
O15484	O15484	CAN5_HUMAN	CAPN5	1338.34			2416.49	10552.7	8818.7	6115.54	2116.34	6158.58
O15498	O15498	YKT6_HUMAN	YKT6				6812.96	7568.54	4067.1	7262.82	2147.43	9695.11
O15511	O15511	ARPC5_HUMAN	ARPC5	5849.04			4371.72		5742.85		5845.58	
O15523	O15523	DDX3Y_HUMAN	DDX3Y	2926.01	2511.1	1982						
O15525	O60675;O15525	MAFG_HUMAN	MAFG				3623.5	2742.13	2258.8		2091.34	
O15533	O15533	TPSN_HUMAN	TAPBP	3668.28			2371.3	1092.53	1019.3	3740.49	2782.56	3029.9
O43143	O43143	DHX15_HUMAN	DHX15	6098.92	6313.98	6435.8	4859.19	23464	12681.7	10372.9	2775.58	8752.82
O43148	O43148	MCES_HUMAN	RNMT					3415.17	3673.78	1664.72		
O43155	O43155	FLRT2_HUMAN	FLRT2	3904.94	1893.78	4508.89						
O43156	O43156	TTI1_HUMAN	TTI1				1267.68	1660.28	1189.92	1251.36	971.353	1483.41
O43169	O43169	CYB5B_HUMAN	CYB5B				1489.97	8098.14	6250.82	16175.1	3034.98	12357.6
O43172	O43172	PRP4_HUMAN	PRPF4	908.551	1382.62	1868.82	1750.72	3051.14	3632.84	3154.51	1374.48	2985.27
O43175	O43175	SERA_HUMAN	PHGDH	26855.1	26694.4	27201.9	27384.2	27624.6	20738.7	17905.5	20277.4	17673.1
O43237	O43237	DC1L2_HUMAN	DYNC1LI2	5337.77	4403.43	9091.24	6399.85	6893.01	4703.13	5773.72	8258.63	5141.96
O43242	O43242	PSMD3_HUMAN	PSMD3	12265.2	15296.5	12412.7	18662.6	18041.2	15827.9	12553	16867.2	14032.4
O43251	O43251;Q9NWB1	RFOX2_HUMAN	RBFOX2				1723.11		685.832	734.779	1898.76	2315.85
O43252	O43252	PAPS1_HUMAN	PAPSS1			1840.49		1653.12	1979.32			
O43264	O43264	ZW10_HUMAN	ZW10				5825.79	4776.96	2575.49	5594.75	5611.04	5172.73
O43290	O43290	SNUT1_HUMAN	SART1				389.821	1412.98	798.826	981.978	457.141	498.151
O43294	O43294	TGFI1_HUMAN	TGFB1I1				1680.28	4418.1	2510.39			
O43301	O43301	HS12A_HUMAN	HSPA12A							4169.83	4518.89	1194.51
O43324	O43324	MCA3_HUMAN	EEF1E1		1659.29	3687.59	5304.24	2822.13	2230.92	2726.58	3868.22	3396.7
O43353	O43353	RIPK2_HUMAN	RIPK2							1429.27	1431.43	2076.89
O43390	O43390	HNRPR_HUMAN	HNRNPR		3338		2940.32	11880.6	3898.8	3405.09		3971.71
O43396	O43396	TXNL1_HUMAN	TXNL1	5411.53		6298.89	1555.87	2211.47	13720.3	2626.83		4235.2
O43399	O43399	TPD54_HUMAN	TPD52L2	9188.54		8739.52	14275.2	5939.43	15568.9	5414.28	4973.61	6695.19
O43427	O43427	FIBP_HUMAN	FIBP							1780.93		1914.46
O43447	O43447	PPIH_HUMAN	PPIH				1749.89	1258.4	1999.24	1360.24		1198.67
O43488	Q8NHP1;O43488	ARK72_HUMAN	AKR7A2			1716.69			2515.86	1746.65	1471.55	2228.94
O43491	O43491	E41L2_HUMAN	EPB41L2	5189.44		5065.13	3350.03	3583.28	4130.15	5301.37	2483.08	5654.66
O43493	O43493	TGON2_HUMAN	TGOLN2				2303.82		635.919		1136.93	
O43502	O43502	RA51C_HUMAN	RAD51C				2529.51			3999.95	7483.3	5460.01
O43542	O43542	XRCC3_HUMAN	XRCC3				1048.35		834.28	2557.68	971.83	2487.95
O43561	O43561	LAT_HUMAN	LAT				1941.24	2691.33	3498.4	7505.79	1858.54	5768.1
O43592	O43592	XPOT_HUMAN	XPOT	1800.72	4920.05	4661.28	7167.46	8953.78	8852.34	6915.87	5603.97	9805.99
O43615	O43615	TIM44_HUMAN	TIMM44			2694.34	6603.41	4328.01	5872.04	7109.2	5667.63	7152.28
O43663	O43663	PRC1_HUMAN	PRC1					2985.16	2138.37	960.875		
O43670	O43670	ZN207_HUMAN	ZNF207				876.172	928.131	1318.43	2241.26	615.663	2089.83
O43681	O43681	GET3_HUMAN	GET3	11058.7	6025.44	9133.85	4458.35		5650.33		6495.83	
O43684	O43684	BUB3_HUMAN	BUB3	4840.72		3566.58	3354.02	6482.33	3876.24	3890.63	2702.35	4427.83
O43707	O43707	ACTN4_HUMAN	ACTN4	21070.6	13234.6	25012.6	38826.6	33311.1	57161.7	37580.2	13721.6	32202.9
O43747	O43747	AP1G1_HUMAN	AP1G1	3949.63	3510.26	3307.29	4765.03	5578.97	5236.7	8532.6	5584.2	7430.38
O43765	O43765	SGTA_HUMAN	SGTA				2252.09	1601.81	2005.62	1912.74		2210.07
O43768	O43768	ENSA_HUMAN	ENSA						1870.93	967.534		874.112
O43776	O43776	SYNC_HUMAN	NARS1	3531.48		6798.85	6486.18	6837.14	15909.4	2933.8	1338.05	3164.56
O43790	O43790	KRT86_HUMAN	KRT86	6742.84		4748.26	6477.45		1156.7	2469.11	4117.86	2217.76
O43795	O43795	MYO1B_HUMAN	MYO1B	27567.7	36685.8	30333.6	22979.1	27406.5	20981.3	81331.7	26118.2	67604.3
O43809	O43809	CPSF5_HUMAN	NUDT21	7012.92	7652.71	6701.72	9683	81458.2	11261	7423.12	8583.36	7381.49
O43815	O43815;Q13033;Q9NRL3	STRN_HUMAN	STRN			1777.88	2069.38	2482.02	3808.94	2390.86		2151.9
O43824	O43824	GTPB6_HUMAN	GTPBP6	2476.07	1921.01	2183.68	1026.41	3282.37	1882.44	2127.74	1110.47	2587.91
O43837	O43837	IDH3B_HUMAN	IDH3B	6536.74	9476.73	4271.27	7151.66	7980.23	5601.73	11715.5	9668.27	9601.76
O43847	O43847	NRDC_HUMAN	NRDC				3491.08	1514.93	3921.01	1740.76		
O43852	O43852	CALU_HUMAN	CALU	15705.5	7550.11	16949.2	17965.6	8165.06	14475.8	4719.59	7774.84	4787.25
O43854	O43854	EDIL3_HUMAN	EDIL3		2252.48	3550.21	6751.96	9253.56	11751.5		6738.18	
O43896	Q12756;O43896;O60333	KIF1C_HUMAN	KIF1C			9236.15						
O43913	O43913	ORC5_HUMAN	ORC5					5379.11	2398.2	3761.33	2535.32	946.475
O43929	O43929	ORC4_HUMAN	ORC4					9181.07	3938.04	4931.71	7496.07	
O60218	O60218	AK1BA_HUMAN	AKR1B10			4040.22						
O60220	O60220	TIM8A_HUMAN	TIMM8A				2929.88		2398.2	2363.32		2152.74
O60231	O60231	DHX16_HUMAN	DHX16					1323.38	843.221	706.645		
O60232	O60232	ZNRD2_HUMAN	ZNRD2				5334.65				4585.28	
O60237	O60237	MYPT2_HUMAN	PPP1R12B					1745.54	1018.99			
O60245	O60245	PCDH7_HUMAN	PCDH7				2282.87	4787.58	5343.93			
O60248	O60248	SOX15_HUMAN	SOX15					731.072				
O60256	O60256	KPRB_HUMAN	PRPSAP2					3443.56	4988.04	2213.14		
O60264	O60264;P28370	SMCA5_HUMAN	SMARCA5			3629.26	3734.11	6729.36	3789.21	2676.73	2641.68	3639.2
O60271	O60271	JIP4_HUMAN	SPAG9				2603.84	2382.29	3453.55	1729.78		1945.46
O60285	O60285	NUAK1_HUMAN	NUAK1	50380		46929.3				5616.05	12708.3	5729.02
O60313	O60313	OPA1_HUMAN	OPA1		2453.12	1754.07	5221.17	4973.4	5245.73	5977.36	4923.58	5707.16
O60315	O60315	ZEB2_HUMAN	ZEB2	2363.06		1581.34						
O60341	O60341	KDM1A_HUMAN	KDM1A					5331.9	3926.81	4698.41	5002.38	3394.28
O60437	O60437	PEPL_HUMAN	PPL	21495.9	10886.2	2853.21		1448.01	657.715	2692.64	1688.3	2150.29
O60476	O60476	MA1A2_HUMAN	MAN1A2				7240.63	1329.56			8817.15	
O60488	O60488	ACSL4_HUMAN	ACSL4	9527.4	9614.64	9827.08	6795.06	6308.35	6572.38	6570.47	4822.45	6084.43
O60493	O60493	SNX3_HUMAN	SNX3			1966.64			2262.73			
O60502	O60502	OGA_HUMAN	OGA			593.911		2013.09	1427.53	1388.69	809.037	2533.69
O60504	O60504	VINEX_HUMAN	SORBS3									
O60506	O60506	HNRPQ_HUMAN	SYNCRIP	9505.59	11838.6	7841.44	7196.71	19784.2	13126.5	9534.64	7289.45	9920.3
O60518	O60518	RNBP6_HUMAN	RANBP6				778.48	9160.2	7311.7	3022.4	1858.59	5518.98
O60547	O60547	GMDS_HUMAN	GMDS			9921.97	3415.65			7386.35	12231.7	8367.56
O60568	O60568	PLOD3_HUMAN	PLOD3				4055.11	4668.38	4815.47	2081.96	5487.53	2268.84
O60573	O60573	IF4E2_HUMAN	EIF4E2				662.284	644.822	560.138	1122.31		
O60609	O60609	GFRA3_HUMAN	GFRA3							5156.39	896.96	6261.15
O60610	O60610	DIAP1_HUMAN	DIAPH1				6642.27	6898.82	6934.24	7576.42	5881.77	7385.93
O60645	O60645	EXOC3_HUMAN	EXOC3	5658.24	7944.96	3903.85	7774.56	5960.97	6878.31	5681.29	4542.56	5881.12
O60656;[illegible]	Q9HAW9;O60656;P22310;P22309;P19224;P35503;P3	UD110_HUMAN;UD11_HUMAN;UD13_HUMA	UGT1A1;UGT1A10;UGT1A3;UGT1A4;UGT1A5;UGT1A6;UGT1A7;UGT1A8;UGT1A9									
O60658	O60658	PDE8A_HUMAN	PDE8A					1139.45	1084.05	2856.67	993.044	3518.25
O60664	O60664	PLIN3_HUMAN	PLIN3	5988.84	2782.85	6294.17	4781.06	6210.43	10487.9	4118.47	1318.73	4329.52
O60678	O60678	ANM3_HUMAN	PRMT3				2244.96	2237.97	4585.53	2124.33	1934.33	2139.73
O60684	O60684	IMA7_HUMAN	KPNA6			2969.1	5035.6		2035.58	3117.17	3421.56	3313.5
O60701	O60701	UGDH_HUMAN	UGDH	9237.86	8027.53	14670.9	4098.05	4002.64	4894.8	5828.44	6066.77	6700.48
O60716	O60716	CTND1_HUMAN	CTNND1	6535.36	6418.93	5667.72	8466.61	12537.8	9757.8	22932.8	11245.1	20792.7
O60732	O60732	MAGC1_HUMAN	MAGEC1				12771.3	10223.6	11676.3			
O60739;P41567	O60739;P41567	EIF1B_HUMAN;EIF1_HUMAN	EIF1;EIF1B				1662.95	1449.97	2793.07	1379.23		
O60749	Q13596;O60749	SNX2_HUMAN	SNX2	4313.9		3577.66	6850.83	3439.85	4756.07	2311.73	2431.77	2825.08
O60762	O60762	DPM1_HUMAN	DPM1	19841	58012	26301.4	9662.15	13914.5	11688	20452.7	15795.9	17067
O60763	O60763	USO1_HUMAN	USO1	5261.58	3335.5	4672.54	5321.65	5624.19	6431.3	3875.4	3872.71	3217.66
O60826	O60826	CCD22_HUMAN	CCDC22	1634.05	1275.28	1093.99		9245.23	4391.55	1263.46	1156.29	940.325
O60828	O60828	PQBP1_HUMAN	PQBP1				687.293	5173.39	6328.25	3466.57		3509.75
O60832	O60832	DKC1_HUMAN	DKC1		2350.83	1364.25	999.132	6221.86	2269.33	5740.06	3581.8	2532.19
O60841	O60841	IF2P_HUMAN	EIF5B	15244.1	25088.4	13659.9	14322	22441.8	15262	15196.4	10162	13414.7
O60869	O60869	EDF1_HUMAN	EDF1				3273.16	1231.1	3583.88		1974.63	1611.26
O60884	O60884	DNJA2_HUMAN	DNAJA2				11058.7	7185.23	7393.12	11665.5	17806	12611.6
O60888	O60888	CUTA_HUMAN	CUTA	15657.5	5545.9	11201		6326.9	26811.5	17700.5		
O60907;Q9BZK7	O60907;Q9BZK7	TBL1R_HUMAN;TBL1X_HUMAN	TBL1X;TBL1XR1			3840.75	967.42	1091.95	1112.26	2094.89		1153.83
O60911	O60911	CATL2_HUMAN	CTSV				2097.67				2345.38	2195.74
O75027	O75027	ABCB7_HUMAN	ABCB7			2576.27	2167.35			2885.78	5253.18	2844.6
O75044	P0DJJ0;O75044;P0DMP2	SRGP2_HUMAN	SRGAP2				964.93		1126.05		1037.39	3084.41
O75083	O75083	WDR1_HUMAN	WDR1	20624.6	16308.2	22133.9	16157.5	21431.6	28422.3	17005	9429.91	11701.4
O75122	O75122	CLAP2_HUMAN	CLASP2		1325.83		7756.97	5114.92	3768.4	2460.89	3794.43	3820.16
O75127	O75127	PTCD1_HUMAN	PTCD1					1361.26		1287.44	3272.77	2686.8
O75131	O75131	CPNE3_HUMAN	CPNE3	7343.21	4752.94	6488.19	5418.57	3185.55	8875.12	5202.62	5225.39	5596.91
O75145	O75145	LIPA3_HUMAN	PPFIA3			3426.13			33279.9	17879.3		13955.2
O75146	O75146	HIP1R_HUMAN	HIP1R				4911.53	5839.06	7564.24	4604.51		5411.72
O75147	O75147	OBSL1_HUMAN	OBSL1			278.682	524.741	435.872	372.121	496.549		628.858
O75150	Q5VTR2;O75150	BRE1B_HUMAN	RNF40				1779.86		2369.69	2697.51		
O75153	O75153	CLU_HUMAN	CLUH				3818.19		2387.51	3884.78	3335.03	4281.57
O75165	O75165	DJC13_HUMAN	DNAJC13	5509.18	6802.85	4108.07	3444.5	4159.6	3391.53	10953.4	5542.56	7345.49
O75175	O75175	CNOT3_HUMAN	CNOT3			1178.82	1717.03	448.627	960.663	1169.1	830.318	895.94
O75179	O75179;Q8IWZ3	ANR17_HUMAN	ANKRD17				4157.79	3738.54	2913.09	3040.23	5368.29	3008.81
O75190	O75190	DNJB6_HUMAN	DNAJB6				6285.85	6061.25	4110.42	2504.84	9767.85	3158.91
O75223	O75223	GGCT_HUMAN	GGCT	15986.4		8372.96	2777.07		4741.22	3283.16	3063.09	3496.86
O75306	O75306	NDUS2_HUMAN	NDUFS2	3411.93		2875.94	3049.08	3274.92	3589.59	7742.41	3510.63	5076.65
O75312	O75312	ZPR1_HUMAN	ZPR1					4375.56	2724.34	1841.12		
O75323	O75323	NIPS2_HUMAN	NIPSNAP2	6978.62		6617.94	5254.77	6842.83	2955.99	6281.92	5202.53	6744.26
O75326	O75326	SEM7A_HUMAN	SEMA7A				1401.36	3225.32	2641.78	1519.67		1099.97
O75340	O75340	PDCD6_HUMAN	PDCD6	3213.34		3743.6	3565.23	5157.15	4277.52	5323.73	3339.74	5376.76
O75342	O75342	LX12B_HUMAN	ALOX12B	20817.3	22403.3	20928.1	6764.29	6809.42	4573.58	5107.91	6082.18	7578.62
O75347	O75347	TBCA_HUMAN	TBCA				2164.13		8986.57	3580.82	2915.73	4329.1
O75348	O75348	VATG1_HUMAN	ATP6V1G1	1160.82			1942.92	1948.9	2426.06	1991.95	1219.15	1427.47
O75351	O75351	VPS4B_HUMAN	VPS4B	3119.63		2489.43	2102.97	1387.45	2840.57	1261.13	394.191	1277.12
O75352	O75352	MPU1_HUMAN	MPDU1				519.436	3185.44	3469.98	3987.46	1719.23	2747.92
O75362	O75362	ZN217_HUMAN	ZNF217	1448.95	823.531	1299	898.84	1323.85	1304.97	647.737		778.156
O75367	O75367	H2AY_HUMAN	MACROH2A1	6574.3	6483.17	3746.61	3114.4	5039.6	4890.77	756.096	2868.11	
O75368	O75368	SH3L1_HUMAN	SH3BGRL				509.066	821.661	2047.86	704.659	421.717	438.151
O75369	O75369	FLNB_HUMAN	FLNB	17470.2	11641.3	16968.8	26910.1	28320.2	27935.1	15186.8	8875.2	13978.3
O75380	O75380	NDUS6_HUMAN	NDUFS6	801.999			238.506	223.895	240.94	1023.18	516.816	587.344
O75381	O75381	PEX14_HUMAN	PEX14							2443.29		2406.22
O75390	O75390	CISY_HUMAN	CS	9025.2		9957.44	24082.9		14682.2	19427.1	22149.1	26097.1
O75394	O75394	RM33_HUMAN	MRPL33	2097.47		3589.1	939.328		1162.04	1148.75	2112.27	1589
O75396	O75396	SC22B_HUMAN	SEC22B	6799.58	9734	4632.11	3123.9	3033.36	4325.84	2258.95	3580.82	2190.79
O75400	O75400	PR40A_HUMAN	PRPF40A					3515.27	1438.97	1099.6		1127.87
O75419	O75419	CDC45_HUMAN	CDC45							1364.12		
O75420	O75420	GGYF1_HUMAN	GIGYF1					7197.38				
O75427	O75427	LRCH4_HUMAN	LRCH4				2867.44	7719.25	5258.19	3480.59		3027.51
O75436	O75436	VP26A_HUMAN	VPS26A	3642.38		4284.06	3342.12	1125.89	2538.61	1669.86	2758.44	1979.98
O75439	O75439	MPPB_HUMAN	PMPCB	6186.88	3786.68	6286.5	6289.21	2887.08	5941.35	4997.1	7579.51	5580.85
O75475	O75475	PSIP1_HUMAN	PSIP1	6081.34				4135.73	3424.09			1824.36
O75477	O75477	ERLN1_HUMAN	ERLIN1		1748.32	2278.66	683.217			1748.44		1981.66
O75489	O75489	NDUS3_HUMAN	NDUFS3	689.961			2107.23	3943.49	3757.04	5209.43	3259.9	5513.34
O75494	O75494	SRS10_HUMAN	SRSF10	1671.59		1958.47		4965.67	1949.5	2110.74	1138.73	1780.72
O75508	O75508	CLD11_HUMAN	CLDN11	1527.21	2134.62	1433.01				2058.28	628.57	1810.96
O75521	O75521	ECI2_HUMAN	ECI2						1172.04	1212.1		1031.72
O75531	O75531	BAF_HUMAN	BANF1	7005.17	20103.2	5137.56	1500.71	10717.2	4011.3	2610	4198.31	6449.46
O75533	O75533	SF3B1_HUMAN	SF3B1	7271.98	10114.8	5278.2	5626.87	18657	8103.48	8665.09	6108.98	8949.92
O75534	O75534	CSDE1_HUMAN	CSDE1	2254.17		3006.61	2264.46	1490.31	1835.07	3762.19	1661.55	3639.16
O75556	O75556	SG2A1_HUMAN	SCGB2A1						798.029		3577.86	
O75569	O75569	PRKRA_HUMAN	PRKRA			18089.4	1297.65	3484.95	1650.77	3236.93	4381.68	2182.56
O75592	O75592	MYCB2_HUMAN	MYCBP2				2334.97			2333.48	4506.13	2416.46
O75600	O75600	KBL_HUMAN	GCAT		787.94		439.518	2008.11	1053.37	1078.34	2565.42	2397.95
O75608	O75608	LYPA1_HUMAN	LYPLA1			3724.11	3367.39	7440.24	9770.77	4780.11	2734.01	5061.32
O75616	O75616	ERAL1_HUMAN	ERAL1	3923.46	7093.01	3427.61	4745.96	3330.38	3446.55	6875.43	5628.19	7094.58
O75635	O75635	SPB7_HUMAN	SERPINB7			4586.81	2918.12	2134.2	2725.4	2071.89	2301.74	3258.99
O75638	O75638	CTAG2_HUMAN	CTAG2			577.904	19913.5	19014.1	17298.1		1843.73	
O75643	O75643	U520_HUMAN	SNRNP200	6959.93		4824.12	7426.42	9794.72	7796.33	8520.61	9841.26	8375.62
O75663	O75663	TIPRL_HUMAN	TIPRL						1173.09			977.302
O75665	O75665	OFD1_HUMAN	OFD1							1408.46		2519.94
O75676	O75676	KS6A4_HUMAN	RPS6KA4		9994.85	8912.44		1533.15	1619.85	5563.3	6853.05	6266.31
O75694	O75694	NU155_HUMAN	NUP155			17647.6	15696.6	10780.7	11449.5	16850.7	16239.4	18828.2

O75695	O75695	XRP2_HUMAN	RP2	9059.18	1829.37	1477.7	13886.9	42124.6	39828.1	81007.5	17446.3	80400.5
O75718	O75718	CRTAP_HUMAN	CRTAP	4571.42			4561.21	14439.8	12945.9	3162.6	1365.41	3453.46
O75746	O75746	S2512_HUMAN	SLC25A12	4131.87				2751.12	1592.18	2440.46		672.836
O75762	O75762	TRPA1_HUMAN	TRPA1	13103.2	19556.8	16005.3						
O75781	O75781	PALM_HUMAN	PALM				3004.67	4033.25	4257.27	3189.75	5083.58	3950.54
O75787	O75787	RENR_HUMAN	ATP6AP2					2813.81	2393.24	3664.89	962.937	1134.6
O75794	O75794	CD123_HUMAN	CDC123	4836.42			6751.68	5093.65	5461.12	4581.93	6004.44	4853.07
O75821	O75821	EIF3G_HUMAN	EIF3G	13739.2			12696.7	15719.6	22006.2	13640.7	6222.48	10592.4
O75822	O75822	EIF3J_HUMAN	EIF3J				8500.48	8936.67	26570.7	9208.7	4529.09	9156.75
O75832	O75832	PSD10_HUMAN	PSMD10	2856.44	1519.98	3165.98	7777.99	5825.73	9847.96	11818.8	9392.86	12313.6
O75843	O75843	AP1G2_HUMAN	AP1G2				5751.3	6848.32	5854.61	7853.39	6322.95	7267.41
O75844	O75844	FACE1_HUMAN	ZMPSTE24					2831.47	2184.26	2983.81		1792.05
O75874	O75874	IDHC_HUMAN	IDH1	6308.13	3326.2	6758.09	4507.11	6005.85	13143.1	4761.11	2070.22	4566.67
O75879	O75879	GATB_HUMAN	GATB				4417.09		4485.05	5264.2	4735.85	6096.95
O75880	O75880	SCO1_HUMAN	SCO1			3655.76	2777.33	3806.97	2640.97	6639.12	2133.84	7212.59
O75891	O75891	AL1L1_HUMAN	ALDH1L1				4092.14	8573.55	7074.11			
O75909	O75909	CCNK_HUMAN	CCNK		1801.69		748.933	1057.13	1939.99	2301.16	405.565	2337.58
O75914	O75914	PAK3_HUMAN	PAK3			8628.35	6641.57					
O75923	O75923	DYSF_HUMAN	DYSF				4309.56	6128.93	5232.61			
O75947	O75947	ATP5H_HUMAN	ATP5PD			1896.7			1408.6	1813.73		2068.81
O75955	O75955	FLOT1_HUMAN	FLOT1	9239.59	7901.29	7201.54	11873.6	13141.8	12022.8	60094.6	22808.5	45075.9
O75962	O75962	TRIO_HUMAN	TRIO	1868.35	1900.56		1172.61	2698.43	3010.46	1626.71		1864.98
O75964	O75964	ATP5L_HUMAN	ATP5MG	4652.86	3667.34	5346.58	1143.12	1419.05	1282.77	2642.32	2283.73	2183.83
O76003	O76003	GLRX3_HUMAN	GLRX3			39889.6	4381.1		8771.72		4174.76	
O76009	O76009	KT33A_HUMAN	KRT33A			3018.3						2763.37
O76011	O76011;A0A140TA62	KRT34_HUMAN	KRT34				4575.49		1783.84	2005.11	1920.13	3180.81
O76013	O76013	KRT36_HUMAN	KRT36				3016.8			3038.26		949.719
O76021	O76021	RL1D1_HUMAN	RSL1D1				284.435	2317.25	1219.63	503.616		684.998
O76024	O76024	WFS1_HUMAN	WFS1	2615.23	2226.68	2116.52				11093.3	3471.06	10363.4
O76031	O76031	CLPX_HUMAN	CLPX	7377.51	6732.99	7900.88	9770.99	6785.76	6153.52	7411.3	12910.1	8984.89
O76070	O76070	SYUG_HUMAN	SNCG				5686.35	1735.65	2042.78			
O76094	O76094	SRP72_HUMAN	SRP72	11115.3	9982.42	7377.06	12562.6	8948.11	11787.7	10152.4	6925.61	10015.6
O94760	O94760	DDAH1_HUMAN	DDAH1			2090.38	5100.52	3814.47	13750.8	2205.78		1854.84
O94766	O94766	B3GA3_HUMAN	B3GAT3						1394.57	2262.63		1652.3
O94776	O94776	MTA2_HUMAN	MTA2	3173.48	3150.3	3373.28	5267.74	5126.94	5725.25	13377.1	5608.68	11732
O94804	O94804	STK10_HUMAN	STK10				1264.99	1875.68	1736.75			1023.71
O94808	O94808	GFPT2_HUMAN	GFPT2					4496.74	3116.99			
O94822	O94822	LTN1_HUMAN	LTN1		742.174	991.359	2656.55	5013.81	4151.63	1725.84	2748.25	1952.36
O94826	O94826	TOM70_HUMAN	TOMM70	1702.92	1743.58	1362.26	1465.38	4332.71	4355.69	3239.69	1258.97	4453.17
O94829	O94829	IPO13_HUMAN	IPO13					776.751	1592.23	2008.37	764.518	2907.73
O94832	O94832	MYO1D_HUMAN	MYO1D						2364.53	27954.2	8208.35	19237
O94842	O94842	TOX4_HUMAN	TOX4				3296.47	1566.3	1808.53	1963.81	5552.84	2578.95
O94851	O94851	MICA2_HUMAN	MICAL2	3329.7			2122.43					
O94874	O94874	UFL1_HUMAN	UFL1			7529.99	2802.22	2490.59	1589.44	2924.59	3140.34	4860.81
O94903	O94903	PLPHP_HUMAN	PLPBP					517.738	1052.35			
O94905	O94905;O75477	ERLN2_HUMAN	ERLIN2	7676.07	11334.7	10593.6	6521.2	12198.1	8073.82	13873	4604.85	11621.8
O94906	O94906	PRP6_HUMAN	PRPF6	8045.67	7115.98	7206.16	12609	8362.61	5649.59	9927.43	18329.4	10261.6
O94915	O94915	FRYL_HUMAN	FRYL				4618.36	2153.05	3364.01	5217.19	5465.86	4650.53
O94925	O94925	GLSK_HUMAN	GLS	5101.88	1686.89	7364.84	9094.91	10259.8	10102	7571.32	11120.5	8916.18
O94927	O94927	HAUS5_HUMAN	HAUS5				1876.15	3455.22	2037.61	3266.98	2389.52	3166.94
O94964	O94964	SOGA1_HUMAN	SOGA1				3565.64	3301.01	2697.62	2811.15		
O94973	O94973	AP2A2_HUMAN	AP2A2				5900.69	1333.91	3114.45	1065.52	1982.16	
O94979	O94979	SC31A_HUMAN	SEC31A	8495.41	6309.88	7843.98	5425.47	7312.62	8630.97	5614.15	6564.83	5222.58
O94992	O94992	HEXI1_HUMAN	HEXIM1				2362.13	2311.46	3780.16	1453.58	880.115	1520.44
O95067	O95067	CCNB2_HUMAN	CCNB2							2544.32	2012.06	2341.47
O95071	O95071	UBR5_HUMAN	UBR5				1252.75		839.864	2560.8	2190.2	2990.55
O95084	O95084	PRS23_HUMAN	PRSS23					7842.32	3944.42			
O95104	Q9UPN6;O95104	SCAF4_HUMAN	SCAF4	5884.62	3017.48	4157.18	4157.39	4097.38	1870.16	3465.39	4167.22	3114.59
O95155	O95155	UBE4B_HUMAN	UBE4B					1919.51	1867.88	1558.13		1404.72
O95163	O95163	ELP1_HUMAN	ELP1		5360.29	1129.46	3985.84	2974.48	3284.45	4360.67	4555.53	4756.61
O95182	O95182	NDUA7_HUMAN	NDUFA7						463.101	696.844		533.827
O95197	O95197	RTN3_HUMAN	RTN3				2822.02	1825.09	2954.15			
O95202	O95202	LETM1_HUMAN	LETM1			1600.85		1859.02	1475.45	2425.74	919.279	2393.12
O95219	O95219	SNX4_HUMAN	SNX4	2024.96	2727.66	2053.6	2222.41	6552.2	3952.11	4763.93	2468.25	2819.93
O95235	O95235	KI20A_HUMAN	KIF20A					1381.45	1133.38	937.088		1239.21
O95239	O95239	KIF4A_HUMAN	KIF4A				2841.27	4415.83	3376.46	4412.64	2464.84	4693.44
O95248	O95248	MTMR5_HUMAN	SBF1				985.161	1580.84		1709.61		2037.42
O95292	O95292	VAPB_HUMAN	VAPB			1939.51	1575.84	1787.27	2359.02	6330.82	2494.27	4937.41
O95297	O95297	MPZL1_HUMAN	MPZL1						10880	7661.8		6773.48
O95336	O95336	6PGL_HUMAN	PGLS	10722.9	2113.17	10492.6	7677.99		1732.34	2418.58	5942.38	3108.11
O95340	O95340	PAPS2_HUMAN	PAPSS2	7976.52		3953.42	8326.69	10653.1	10175.6			
O95347	O95347	SMC2_HUMAN	SMC2	4232.83	3090.74	4474.83	6689.31	6314.62	4968.39	6371.26	7816.89	6673.98
O95352	O95352	ATG7_HUMAN	ATG7		2425.65		1354.67	2808.55	3505.3	1003.79		
O95363	O95363	SYFM_HUMAN	FARS2						1259.15	2485.95		2556.61
O95373	O95373	IPO7_HUMAN	IPO7	6284.64	5714.83	8298.63	19811.7	23420.7	22435	13447.3	20941.7	14331.6
O95396	O95396	MOCS3_HUMAN	MOCS3				2883.79			896.314	2225.56	651.331
O95425	O95425	SVIL_HUMAN	SVIL	6319.79	6316.9	5501.21	9579.16	49859	40074.6	9790.87	2616.76	7714.12
O95433	O95433	AHSA1_HUMAN	AHSA1			4535.42	6005.18	5644.76	11688.3	11519	5717.58	10816.3
O95466	O95466	FMNL1_HUMAN	FMNL1				3268.69	14122	10748.1	13520.8	2885.8	9361.08
O95470	O95470	SGPL1_HUMAN	SGPL1	5529.81	12203.2	4950.24	8821.5	9094.46	8053.41	12166.3	13391.4	9298.56
O95486	O95486	SC24A_HUMAN	SEC24A	3015.34	1326.55	1813.73	2513.96	4355.78	1903.72	1809	570.105	1738.11
O95487	O95487	SC24B_HUMAN	SEC24B				1087.76	1903.22	1493.43	929.498		859.655
O95571	O95571	ETHE1_HUMAN	ETHE1	5057.42	13177.7	4503.96	6511.48		4132.25	2932.9	2079.24	5319.3
O95573	O95573	ACSL3_HUMAN	ACSL3	27525.7	25758.4	24040.2	31942.4	25720.9	18362.1	14995.4	26385.2	14967.6
O95613	O95613	PCNT_HUMAN	PCNT	7982.51	8367.88	8521.98		3353.2	2216.44	2008.92		
O95619	O95619	YETS4_HUMAN	YEATS4							1160.38		1250.01
O95630	O95630	STABP_HUMAN	STAMBP	2714.61	1213.65	2914.16	1232.6		3099.31	2029.14		2058.52
O95671	O95671	ASML_HUMAN	ASMTL							2191.09	2022.71	2640.02
O95678	O95678	K2C75_HUMAN	KRT75	3465.61	3323.11		2685.62			1873.7		2609.64
O95714	O95714	HERC2_HUMAN	HERC2	922.376			1665.37	2796.95	1488.86	1534.23	1909.64	2197.44
O95716;P20337;Q96E17	P20337;O95716;Q96E17	RAB3B_HUMAN;RAB3C_HUMAN;RAB3D_HU	RAB3B;RAB3C;RAB3D									
O95721	O95721	SNP29_HUMAN	SNAP29					689.895	1048.52	779.574		
O95747	O95747	OXSR1_HUMAN	OXSR1	8391.57	3707.55	9388.15	13397.9	6135.34	7679.82	8342.72	10279.1	5348.75
O95758	O95758	PTBP3_HUMAN	PTBP3	5389.39		4174.07	4477.27	10129.7	3601	2978.73	4042.31	3402.16
O95782	O95782;O94973	AP2A1_HUMAN	AP2A1	6759.33		4540.36	6519.31	10343.2	10883.8	5759.81	3146.29	4689.2
O95785	O95785	WIZ_HUMAN	WIZ				812.466	4348.63				
O95786	O95786	RIGI_HUMAN	RIGI	1083.72			3867.32	2925	3428.8			
O95793	O95793	STAU1_HUMAN	STAU1	5234.02	9383.26	5830.78	12786.9	18156.8	10658.2	7107.98	9165.55	8222.81
O95801	O95801	TTC4_HUMAN	TTC4				5826.58	10129.1	10907.9	6873.4		5826.39
O95810	O95810	CAVN2_HUMAN	CAVIN2		545.995					3031.32		
O95816	O95816	BAG2_HUMAN	BAG2	5242.96	4870.62	4339.12	4500.59	11292.1	10825	4991.67	2494.57	5736.54
O95817	O95817	BAG3_HUMAN	BAG3				2917.42	2188.09	3486.71			
O95819	O95819	M4K4_HUMAN	MAP4K4			1029.79	2657.75	2045.31	1255.39	2846.72	1127.31	2033.52
O95822	O95822	DCMC_HUMAN	MLYCD	5411.09		4303.1	6962.4	5434.11	5453.19	5288.33	5418.31	5231.39
O95831	O95831	AIFM1_HUMAN	AIFM1	5297.85	5121.37	4961.15	6136.94	6034.09	2672.61	8647.02	24666.1	8120.27
O95837	O95837	GNA14_HUMAN	GNA14					2655.3	3082.93	4260.21		5130.15
O95864	O95864	FADS2_HUMAN	FADS2									
O95865	O95865	DDAH2_HUMAN	DDAH2			1752.83		285.383	1442.94	953.307	587.972	776.748
O95881	O95881	TXD12_HUMAN	TXNDC12					996.211	5565.04	1566.11		
O95968	O95968	SG1D1_HUMAN	SCGB1D1								13255	
O95994	O95994	AGR2_HUMAN	AGR2									
O96000	O96000	NDUBA_HUMAN	NDUFB10		1346.29		2136.9					
O96005	O96005	CLPT1_HUMAN	CLPTM1	7857.59			1270.5	3769.7	3026.05	3547.68		1397.03
O96008	O96008	TOM40_HUMAN	TOMM40				3169.09		872.299	4225.52	1853.83	4164.36
O96013	O96013	PAK4_HUMAN	PAK4				2252	1095.48	2407.86	2405.47	2664.36	2380.45
O96019	O96019	ACL6A_HUMAN	ACTL6A				3839.07		1643.72	3070.99	4354.41	2127.95
P00338	P00338	LDHA_HUMAN	LDHA	19964.3	8443.23	23599	16917.2	7948.65	23106.9	13341.6	10683.6	13883.4
P00352	P00352	AL1A1_HUMAN	ALDH1A1	7309.34		6326.17	2654.71			1857.89		2626.76
P00367	P00367;P49448	DHE3_HUMAN	GLUD1	171196	545082	20776.9	13716.3	30210.2	16005.4	14017.1	12099.2	17354.5
P00387	P00387	NB5R3_HUMAN	CYB5R3	6725.85	6022.37	12419.7	4258.69	7129.07	6612.73	5546.15	2998.38	4787.62
P00390	P00390	GSHR_HUMAN	GSR	5856.24		5882.57	4765.66	3555.23	3459.99	2787.48	3376.16	3145.06
P00403	P00403	COX2_HUMAN	MT-CO2	8666.82		10245.3	3975.27	8131.36	8422.57	6189.54	5486.94	11128.2
P00441	P00441	SODC_HUMAN	SOD1	10095		15325.1	13329.2	7822.8	26806.1	6528.74	8729.98	14309.3
P00491	P00491	PNPH_HUMAN	PNP	6536.5	7159.92	6132.16	4507.37	3646.85	8317.19	4195.35	2437.99	4844.35
P00492	P00492	HPRT_HUMAN	HPRT1	2768.66		2768.74	3808.61	1940.86	5254.97	5352.43	3148.58	4601.82
P00505	P00505	AATM_HUMAN	GOT2	8824.02		7245.65	8094.2	25617.2	6447.93	11892.3	9843.6	11614.6
P00533	P00533	EGFR_HUMAN	EGFR	7508.08	8189.72	6851.71	5529.56	7330.28	8296.36			
P00558	P00558	PGK1_HUMAN	PGK 1.00	16453.9	10576.7	23062.8	33547.5	36381.5	80384.5	34485.1	14111.3	22880
P00568	P00568	KAD1_HUMAN	AK1				7054.36		3196.49	809.698		
P00750	P00750	TPA_HUMAN	PLAT	12428	12505.8	13756.8						
P00966	P00966	ASSY_HUMAN	ASS1	2130.1		2547.24				2332.92	6008.43	4849.41
P01023	P01023	A2MG_HUMAN	A2M	8580.14	4223.54	10803.6	1427.09		2683.75	8724.03	1651.52	5558.18
P01024	P01024	CO3_HUMAN	C3			4712.74				438.05	804.905	
P01036	P01036;P09228;P01037	CYTS_HUMAN	CST4	23997.1	4242.12	5459.29	3448.94	1575.55	2421.26	1652.34	5627.81	2532.48
P01040	P01040	CYTA_HUMAN	CSTA	135845	58270.3	192282	84508.1	31787.7	55281.4	28068.3	82188	62284.8
P01112	P01111;P01112	RASH_HUMAN	HRAS					4420.31	3665.69	7668.42		7918.72
P01116	P01116	RASK_HUMAN	KRAS	9720.52		8627.49	13032.9	30209.7	26476.3	59815.8	21477.5	129322
P01591	P01591	IGJ_HUMAN	JCHAIN	6877.84	4538.06	15915.2					3186.55	
P01833	P01833	PIGR_HUMAN	PIGR	1931.22		5560.19				1206.93	1629.68	1151.89
P01834	P01834	IGKC_HUMAN	IGKC	12207.1	6724.47	39831.4	6289.88	12461.6	11031.4	7582.88	6615.91	9670.66
P01857	P01859;P01857	IGHG1_HUMAN	IGHG1	16738.3	14254.9	57718.7	11891.9		6884.34	6189.81	12969.9	9713.25
P01859;P01861	P01859;P01861	IGHG2_HUMAN;IGHG4_HUMAN	IGHG2;IGHG4	22068.6	9861.1	98241.9	13208.2			9232.88	12782.2	
P01876	P01876;P01877	IGHA1_HUMAN	IGHA1	25780.5	9775.02	49695.4	10868.1	3325.76	2485.17	6136.87	15900.5	10342.3
P01877	P01877	IGHA2_HUMAN	IGHA2								1541.11	
P01889	P01889	HLAB_HUMAN	HLA-B	2566.54	7039.28	2582.26			523.38	753.957		641.634
P01893	P01893	HLAH_HUMAN	HLA-H									
P02144	P02144	MYG_HUMAN	MB									
P02452	P02452	CO1A1_HUMAN	COL1A1				7672.89				5398.03	
P02533	P02533	K1C14_HUMAN	KRT14	490084	501499	851711	296101	137985	185851	246508	546638	447118
P02538	P02538;O95678;P48668	K2C6A_HUMAN	KRT6A	195361	160816	383792	119137	39130	46141.8	70794.8	232837	167403
P02545	P02545	LMNA_HUMAN	LMNA	37267.3	32605.4	31769.1	18256.9	18703.9	25091.5	19974.1	9273.9	18131
P02585	P02585	TNNC2_HUMAN	TNNC2									
P02689;P15090	P02689;P15090	FABP4_HUMAN;MYP2_HUMAN	FABP4;PMP2									
P02751	P02751	FINC_HUMAN	FN1	1052.19		2391.13		2893.82	1503.67			
P02765	P02765	FETUA_HUMAN	AHSG	3041.58	1880.4	3003.48				2171.06	571.637	1713.11
P02768	P02768	ALBU_HUMAN	ALB	250511	118706	547603	333896	128172	133485	156996	375895	261168
P02786	P02786	TFR1_HUMAN	TFRC	23429.7	20970	18347	17157.8	12948.4	14377.9	34819	20574.5	25433.3
P02787	P02787	TRFE_HUMAN	TF			4076.9			20676.6			
P02788	P02788	TRFL_HUMAN	LTF	8399.56	7873.39	16776.9	5860.06	2680.11	3694.46	3476.52	10006.6	4465.87
P02792	P02792	FRIL_HUMAN	FTL	15452.4	18293.1	14190.9	8358.18	15483.2	21388.1	3388.51	4297.87	4722.35
P02794	P02794	FRIH_HUMAN	FTH1	4686.21	2849.39	4732.74	5696.6	2204.77	4102.74	5892.57	4910.63	6139.41
P03956	P03956	MMP1_HUMAN	MMP1							1683.2		
P03973	P03973	SLPI_HUMAN	SLPI	4834.32	2385.05	18837.7	2342.26		1854.2	4596.95	1695.36	2201.11
P04004	P04004	VTNC_HUMAN	VTN						1111.62	2922.71		
P04040	P04040	CATA_HUMAN	CAT	12882.6	9387.11	20065.6	6791.82	7742	13134.6	8953.04	5844.75	10394.2
P04049	P04049	RAF1_HUMAN	RAF1	3769.94			2175.85		1216.57	5557.32	2432.88	5296.6
P04062	P04062	GBA1_HUMAN	GBA1	3235.53		2311.36	1495.69	2539.05	1977.48	2064.02	1058.39	1334.79
P04075	P04075	ALDOA_HUMAN	ALDOA	36473.6	14352.9	52889.9	37868.7	27676.8	68000.9	38061.4	16427.6	39489.1
P04080	P04080	CYTB_HUMAN	CSTB	20380.2	9300.69	12005.9	5536.76	4218.01	7454.11	6661.76	6988.24	4567.85
P04083	P04083	ANXA1_HUMAN	ANXA1	37864.7	16030	43139	26601.5	26417.4	99361.6	14752.4	6222.02	15881.2
P04114	P04114	APOB_HUMAN	APOB		2045.66	2548.76	3336.49	2182	2456.64	798.008		
P04150	P04150	GCR_HUMAN	NR3C1			1605.82	2613.77	1219.21	831.847			
P04156	P04156	PRIO_HUMAN	PRNP				11933.9	26053	13431.5	18736.4	12299.1	21044.9
P04179	P04179	SODM_HUMAN	SOD2	26340	13371.8	54970.9	9455.88		15410.3	18271.1	10587.2	17522.8
P04181	P04181	OAT_HUMAN	OAT	30242.7	28715.1	31212	12641.7	19439	13278.1	105251	123609	105524
P04183	P04183	KITH_HUMAN	TK1			5824.57	5884.52	4284.75	3546.6	5677.07	11041.6	9121.25
P04216	P04216	THY1_HUMAN	THY1	7517.86	8653.46	8778.35	2268.91		1102.87			
P04259	P04259;P48668	K2C6B_HUMAN	KRT6B	17236.8	10378.6	35461.5	17437.2			2263.49	35151.3	10057.8
P04264	P04264	K2C1_HUMAN	KRT1	##########	1.2098E+06	2.20228E+06	1.13055E+06	288127	508459	525788	1.11694E+06	1.17794E+06
P04279	P04279	SEMG1_HUMAN	SEMG1	3840.22		3234.77	3740.4	1257.33	1046.77	676.271	1629.67	717.991
P04350	P04350	TBB4A_HUMAN	TUBB4A	4795.1	9033.25	4119.9	3407.17	26299.9	12846.9	25240.6	14206	6971.8
P04406	P04406	G3P_HUMAN	GAPDH	132935	90889.9	160204	94042.1	91400.8	259847	207111	84557.2	206854
P04439	P01889;P04439	HLAA_HUMAN	HLA-A	14371.4	12675.7	12682.5	3851.91	13077.3	10308.9	33727.3	10172.8	25464.4
P04632	P04632	CPNS1_HUMAN	CAPNS1				4229.99	1142.19	1443.6	3051.47	4130.56	2858.07
P04637	P04637	P53_HUMAN	TP53				1055.05		1452.77	3048.78	3293.59	2992.34
P04792	P04792	HSPB1_HUMAN	HSPB1	125705	135137	133001	40637.3	34845.4	39751	56862.9	57697	70838.4
P04818	P04818	TYSY_HUMAN	TYMS				475.771		1442.32	550.181		754.923
P04843	P04843	RPN1_HUMAN	RPN1	19612.3	18957	12259.5	13947.7	17763	12477.8	20320.9	15904.6	18366.1
P04844	P04844	RPN2_HUMAN	RPN2	6395.89	7822.19	4776.03	10320.8	8392.96	8432.25	9590.99	10972.1	16355.5
P04899	P04899	GNAI2_HUMAN	GNAI2	9331.58	11292.2	11933.2	57119.8	101203	112167	328533	60329.3	315582
P04908;Q7L7L0;Q93077	P04908;Q7L7L0;Q93077	H2A1B_HUMAN;H2A1C_HUMAN;H2A3_HUM	H2AC25;H2AC6;H2AC8	21792.1	29300.5	16150.8	8757.53	30776.7	15389.4	4171.67	4860.38	4568.47
P04920	P04920	B3A2_HUMAN	SLC4A2					652.815	381.571	3461.07		2201.96
P05023	P05023	AT1A1_HUMAN	ATP1A1	16645.5	11771.9	15403.4	13567.2	9211.54	10595.4	33933.5	25520.9	26205.7
P05026	P05026	AT1B1_HUMAN	ATP1B1	9544.76	3868.71	8865.9	13528.4	12148.8	4800.76	24852	16741.8	17318
P05089	P05089	ARGI1_HUMAN	ARG1	49963.4	32074.5	68939.6	28675.7	9525.17	17035.5	21002.3	22343	25246.8
P05090	P05090	APOD_HUMAN	APOD	10868.6	5526.2	12918.4	15952		3680.95	3132.84	8760.14	5901.97
P05091	P05091	ALDH2_HUMAN	ALDH2	12552.1	9445.28	14513.6	789.709		1114.98	21734.9	16843.2	23499.3
P05109	P05109	S10A8_HUMAN	S100A8	99232.9	80410.2	196783	84583.7	13486.1	33704	38148.2	169891	83304.7
P05114	P05114	HMGN1_HUMAN	HMGN1	683.504			2134.34	12034.9	8262.08	1804.97	1063.5	1305.65
P05141	P12236;P05141;P12235;Q9H0C2	ADT2_HUMAN	SLC25A5	30024.5	36838.8	26851.4	14089.3	26161.5	19996	94704.3	59783.6	67819.1
P05161	P05161	ISG15_HUMAN	ISG15	18952.8	13334.9	13799.6	16780.8	7013.1	9600.87		4076.45	
P05164	P05164	PERM_HUMAN	MPO			6636.62	1916.36				2338.48	
P05165	P05165	PCCA_HUMAN	PCCA	610856	507514	775299	571017	103928	156117	370215	993775	578259
P05166	P05166	PCCB_HUMAN	PCCB	433820	261783	596294	389290	54202.4	98658.5	317655	720735	502452
P05187	P05187;P09923	PPB1_HUMAN	ALPP					2851.42		7668.19	4158.68	6569.61
P05198	P05198	IF2A_HUMAN	EIF2S1	12061	7780.9	7112.58	9137.17	14946.7	14433.9	8728.98	5167.26	6958.41
P05204	O00479;P05204	HMGN2_HUMAN	HMGN2	2487.62	2472.97	2314.49	2060.87	1639.83	1271.06	998.25	1917.32	1455.37
P05362	P05362	ICAM1_HUMAN	ICAM1	9844.67	6128.71	9042.37	7071.27	6905.77	4508.1	1683.5		
P05386	P05386	RLA1_HUMAN	RPLP1	9858.75		12242.5	9192.33	291875	132840	81334.5	32643.5	107447
P05387	P05387	RLA2_HUMAN	RPLP2	40293.3	43034.5	37047.4	42148.7	56121	46934.6	45012.7	52582.5	66833.5
P05388	P05388	RLA0_HUMAN	RPLP0	44581.3	49338.8	38363.8	83331	58096	81636.5	79966.2	84333.8	48933.3
P05412	P05412	JUN_HUMAN	JUN	3036.41							1327.87	
P05423	P05423	RPC4_HUMAN	POLR3D	10447	13744.4		4309.35	5984.2	3940.05		2582.71	3193.24
P05455	P05455	LA_HUMAN	SSB	2245.11	1792.99	3824.44	4591.2	6585.16	8753.95	8428.13	4291.79	8090.52
P05549	P05549	AP2A_HUMAN	TFAP2A									
P05556	P05556	ITB1_HUMAN	ITGB1	9240.89	5963.74	10302.3	9974.24	40238.7	45549.3	18962	1880.05	9724.76
P05783	P05783	K1C18_HUMAN	KRT18	220450	85760.6	181543	622476	2.19896E+06	1.55022E+06	1.32455E+06	638500	1.25601E+06
P05787	P05787	K2C8_HUMAN	KRT8	591216	338146	486908	988039	3.62591E+06	2.47228E+06	2.68059E+06	1.20708E+06	2.54681E+06
P05976	P05976;P08590	MYL1_HUMAN	MYL1								1343.71	
P05997	P05997	CO5A2_HUMAN	COL5A2		2150.03		1504.26	8678.84	2207.67			
P06132	P06132	DCUP_HUMAN	UROD	3148.6		5599.29	1175.46	4480.24	12852.6	2946.68		4945.67
P06239	P06239	LCK_HUMAN	LCK				5385.3	730.002	461.761	138497	18148.2	109323
P06241	P06241	FYN_HUMAN	FYN				2671.41	4492.67	4979.8	33723.2	8551.62	25661.6
P06396	P06396	GELS_HUMAN	GSN	8369.03	3016.2	9352.35	11643.8	8967.09	10144.4	3820.51	2436.54	4217.38
P06400	P06400	RB_HUMAN	RB1				6634.8	4820.99	4008.01	5415.4	8067.02	5717.11
P06454	P06454	PTMA_HUMAN	PTMA						2320.5	5145.41	8898.78	5610.76
P06493	P06493	CDK1_HUMAN	CDK1	8455.9	6352.29	8963.01	16922.5	10567	17718.6	15911.2	27855.4	17672.3
P06576	P06576	ATPB_HUMAN	ATP5F1B	37031.6	35972.7	32314.9	21657.2	16706.5	13408.4	40984.3	46181.4	35637.5
P06702	P06702	S10A9_HUMAN	S100A9	30388.2	29189.4	54874.5	23680.8	9447.68	21298.7	14897.9	47682.3	32805
P06703	P06703	S10A6_HUMAN	S100A6	10397.4	2647.42	12682.6	3337.92		2173.04		6388.24	3666.9
P06730	P06730	IF4E_HUMAN	EIF4E	2517.44	1546.57	3361.47	3549.32	3352.8	6144.23	2689.02		1954.22
P06732	P06732	KCRM_HUMAN	CKM									
P06733	P06733	ENOA_HUMAN	ENO1	73002.9	22233.4	80135.3	37111.9	36585	94884.6	71034.3	31810.3	65020.4
P06737	P06737	PYGL_HUMAN	PYGL			4409.63	5217.08	8987.1	11494.1	10026.4	3938.37	10200.4
P06744	P06744	G6PI_HUMAN	GPI	20190.6	11466.2	19339.4	13506.9	11261.6	16328.5	14671.6	13773.4	13015.5
P06746	P06746	DPOLB_HUMAN	POLB	12425.6	21722.7		3915.86	8380.04	2619.45			
P06748	P06748	NPM_HUMAN	NPM1	11119.4	6481.69	10156.4	9900.22	16919.5	18175.1	14670.7	7687.28	15052.1
P06753	P06753	TPM3_HUMAN	TPM3	15928.4	17054.2	15618.6	24456.3	82391.9	64575.5	29439.2	13791.1	23769.9
P06756	P06756	ITAV_HUMAN	ITGAV		5430.44		3014.02	3698.8	2576.22	1931.66		2487.22
P06858	P06858	LIPL_HUMAN	LPL									
P07099	P07099	HYEP_HUMAN	EPHX1	3031.9	4412.35	3763.68		3077.61	2330.43	3115.06		4296.62
P07108	P07108	ACBP_HUMAN	DBI	6339.3	1920.69	6642.81	2242.06	2299.59	7193.34	2431.76	3822.18	4300.91
P07195	P07195;Q6ZMR3;P07864;P00338	LDHB_HUMAN	LDHB	29130.1	13506.1	32603.9	35129.3	9847.07	41845.8	35247.2	24817.9	33877.1
P07237	P07237	PDIA1_HUMAN	P4HB	24045.5	13645.7	24046.3	15521.8	16218.3	42016.8	19601.1	10473.9	16588.8
P07305	P07305	H10_HUMAN	H1-0					971.043	214.807	359.744		143.058
P07339	P07339	CATD_HUMAN	CTSD	16414	22751.6	28292.8	14377.9	7831.44	22491.5	17115.6	14044.8	11133
P07355	P07355;A6NMY6	ANXA2_HUMAN	ANXA2	44109.6	29673.6	60145.3	44719.6	49342.2	126546	46869	25368.5	41473.1
P07384	P07384	CAN1_HUMAN	CAPN1	17416.6	15506.1	17112.8	10608.7	6950.03	6645.83	12924.8	12340.3	14893.8
P07437	P07437;P04350;P68371	TBB5_HUMAN	TUBB	647718	601306	649565	839933	542909	490974	1.03872E+06	1.21602E+06	940640
P07477	Q9BYE2;P07477	TRY1_HUMAN	PRSS1	##########	##########	3.33285E+06	2.36378E+06	381621	463825	1.20685E+06	2.07526E+06	1.30865E+06
P07602	P07602	SAP_HUMAN	PSAP	2547.91		2568.39	1283.89	6715.45	13553	3632.19		
P07686	P07686	HEXB_HUMAN	HEXB	3304.09	1079.02	4025.27	1687.59	2224.77	4082.21	1532.47		1611.48
P07737	P07737	PROF1_HUMAN	PFN1	36245.6	17974.5	31711.6	66791.9	16580.2	61148.8	47782.4	45352.4	25884.7
P07741	P07741	APT_HUMAN	APRT	3726.7		9544.71	15678	22393.8	34342.7	7252.52	6208.36	6774.95
P07814	P07814	SYEP_HUMAN	EPRS1	8268.75	8168.91	6867.13	13036.9	9930.25	11908.1	9238.55	9261.7	9650.35
P07858	P07858	CATB_HUMAN	CTSB	8541.58	12373.1	9778.63	5671.89	6749.47	6112.83	4533.81	16022.2	5044.24
P07900	P07900	HS90A_HUMAN	HSP90AA1	14918.4	13838.7	16207.5	38400.4	36826.4	66766.4	54771	17130.4	52313.6
P07910	P07910	HNRPC_HUMAN	HNRNPC	61429.3	20279.2	13035.5	41792.9	29700.3	9744.67	13668.1	18612	11549.7
P07942	P07942	LAMB1_HUMAN	LAMB1				1461.84	1759.5	2084.53			
P07947	P12931;P07947;P06241	YES_HUMAN	YES1				4741.01	15488.4	9991.13	22676.3	12771.4	20054.7
P07948	P07948	LYN_HUMAN	LYN			20455.7	7497.49	15340.6	12288.4	45006.3	15460.3	40028.8
P07951	P07951	TPM2_HUMAN	TPM2	7443.2		6953.59	12750.6	56810.2	23665.9	8497.54	4972.6	5011.08
P07954	P07954	FUMH_HUMAN	FH	3562.63		2934.48	1373.71	1267.85	2557.72	1736.22	3190.92	2183.2
P07996	P07996	TSP1_HUMAN	THBS1									
P08047	P08047	SP1_HUMAN	SP1				594.337				1073.77	158.543
P08133	P08133	ANXA6_HUMAN	ANXA6	14366.2	7538.11	15712.6	8659.73	8629.47	22188.5			
P08134	P08134	RHOC_HUMAN	RHOC	7679.25	6083.31	9532.59	5855.76	2176.94	5313.83	3282.28	2607.56	3499.67
P08174	P08174	DAF_HUMAN	CD55			2100.76	12417.6	11158.2	8750.57	5323.2		3617.46
P08195	P08195	4F2_HUMAN	SLC3A2	8603.08	10331.3	8653.88	21015.8	23564.4	29172.8	68309.3	19146.2	51962.2
P08236	P08236	BGLR_HUMAN	GUSB					3084.96	1900.85	4146.6	2674.61	4212.11
P08237	P08237	PFKAM_HUMAN	PFKM	10822	14296.9	14248.5	18192.1	23344.4	19418.5	20903.1	17082.3	22353.8
P08238	P08238	HS90B_HUMAN	HSP90AB1	49764.4	44657.8	67838.7	170473	153728	293133	161382	70802.1	166202
P08240	P08240	SRPRA_HUMAN	SRPRA	4314.85	4654.46	2765.49	4990.89	3968.57	3837.11	1729.83	2189.57	2157.77
P08243	P08243	ASNS_HUMAN	ASNS	6911.04	5390.96	5172.96	6978.32	9098.55	10193.1	4239.86	4686.01	5360.41
P08311	P08311	CATG_HUMAN	CTSG			15421.4			2413.95	1765.44	2605.82	
P08397	P08397	HEM3_HUMAN	HMBS						1453.78			
P08473	P08473	NEP_HUMAN	MME	2973.64	5024.44	7705.07	6824.66	11540.5	9492.43	15480.7	4540.38	12447.4
P08559	P08559;P29803	ODPA_HUMAN	PDHA1	3647.79	2270.26	3206.86	9937.88	726.181	5695.97	3237.54	4289.01	3737.56
P08571	P08571	CD14_HUMAN	CD14							4949.22	1143.54	7351.53
P08574	P08574	CY1_HUMAN	CYC1	2939.18	5580.75	3238.35		2789.6	2462.53	4315.15	1442.88	3827.41
P08582	P08582	TRFM_HUMAN	MELTF				6227	7025.01	5621.31	13722.1	8273.92	13329.3
P08621	P08621	RU17_HUMAN	SNRNP70	6002.83	8444.16	5045	5929.18	19861.5	10150.5	11462.9	8325.59	15949.8
P08648	P08648	ITA5_HUMAN	ITGA5					2629.67	1799.62			
P08651	P08651	NFIC_HUMAN	NFIC	56263.1	63785.6	7454.79	3401.13	7940.74	3161.98	5927.85	1739.21	
P08670	P08670	VIME_HUMAN	VIM	1.4686E+07	##########	1.21193E+07	274084	725499	444536	1.15979E+06	233381	1.11681E+06
P08708	P08708	RS17_HUMAN	RPS17	26956.6	38557.1	30762.3	29680	56434.6	31116.8	26978.8	23273.3	31252.2
P08727	P08727	K1C19_HUMAN	KRT19	62972.6	87113.1	62549.8	46927.4	41712.1	23211.4	2.22466E+06	935818	2.00807E+06
P08729	P08729	K2C7_HUMAN	KRT7	107374	47272.9	59278.7	40492.7	11311.4	14746.6	15319.3	32841.1	12979.2
P08754	P08754	GNAI3_HUMAN	GNAI3		5390.45		42627.5	36803.1	32452.3	109524	36329.4	126667
P08758	P08758	ANXA5_HUMAN	ANXA5	31681.1	14611.8	37609.3	13647.7	11154	57211.3	8511.75	4568.97	7845.97
P08779	P08779	K1C16_HUMAN	KRT16	94464.1	87815.6	175613	44214.4	31479.3	30306.2	43595.7	95892.9	82821.6
P08865	A0A8I5KQE6;P08865	RSSA_HUMAN	RPSA	60608.7	68882	52986.1	31440	143908	67002.6	66456.5	32932.1	66498.1
P09012	P09012	SNRPA_HUMAN	SNRPA		8472.42		5727.2		3058.62	2328.57		5087.63
P09104	P09104	ENOG_HUMAN	ENO2			1875.16		612.647	2638.07	944.014		1198.13
P09110	P09110	THIK_HUMAN	ACAA1					837.261	1073.62	1356.17		2800.73
P09132	P09132	SRP19_HUMAN	SRP19			1926.83	3241.43		2933.04	2070.09		3100.63
P09211	P09211	GSTP1_HUMAN	GSTP1	6181.38	3730.32	5820.9	4831.16	5277.11	16720.6	3489.45	5919.64	5928.23
P09327	P09327	VILI_HUMAN	VIL1							9828.88	5444.17	8050.22
P09382	P09382	LEG1_HUMAN	LGALS1	86670.7	35506.9	58814.6	54125.6	70966.7	121350	859501	167864	474050
P09429	P09429;P26583	HMGB1_HUMAN	HMGB1	18409.1	22819.9	16118.2	18298.3	16152	20537.7	15540	14828.4	15351.9
P09471	P09471	GNAO_HUMAN	GNAO1							1013.01		971.695
P09486	P09486	SPRC_HUMAN	SPARC	3102.07		4584.32	13056.3	9233.77	14334.9			
P09493	P09493	TPM1_HUMAN	TPM1	48050.8	38639.8	35261.3	105637	483286	253093	117622	45817.4	72277.5
P09496	P09496	CLCA_HUMAN	CLTA	9913.16			15516.9	15707.9	22799.5	14320.1	8516.06	12124.9
P09497	P09497	CLCB_HUMAN	CLTB				5652.98			3533.65	5433.85	
P09525	P09525	ANXA4_HUMAN	ANXA4			1541.08			2290.69	2144.61		3728.06
P09543	P09543	CN37_HUMAN	CNP	13670.6	16196.7	12798.3	8512.32	8139.25	7512.67	7337.7	6642.41	7782.08
P09601	P09601	HMOX1_HUMAN	HMOX1	3717.91	3005.92	4386.88	1131.19	2728.44	3257.9	1863.02		1697.48
P09619	P09619	PGFRB_HUMAN	PDGFRB						5730.41			
P09622	P09622	DLDH_HUMAN	DLD	16476	11181.4	14198.8	19330.4	12262.6	14032.5	15238.6	14268.8	16105.9
P09651	A0A2R8Y4L2;P09651	ROA1_HUMAN	HNRNPA1	5914.83	3945.55	6421.96	6538.09	70158.9	24596.4	12262.8	6034.71	12373.3
P09661	P09661	RU2A_HUMAN	SNRPA1			3894.89	2824.25	6754.57	6826.62	6985.79	3101.86	7466.19
P09874	P09874	PARP1_HUMAN	PARP1	48531.6	83890	11036.8	71391.2	135026	27195	15148.2	25084.4	17263.5
P09913	P09913	IFIT2_HUMAN	IFIT2				2291.02	2083.23	2067.68			
P09914	P09914	IFIT1_HUMAN	IFIT1	11793.3	11789.6	9243.51	6533	9683.41	10198.8			1276.41
P09936	P09936	UCHL1_HUMAN	UCHL1	3725.97		15678.7						
P09960	P09960	LKHA4_HUMAN	LTA4H			2093.27	1957.76	1509.21	2903.23	2086.32	1482.44	2521.75
P09972	P09972	ALDOC_HUMAN	ALDOC				928.566	2032.89	6625.86	3378.6		3958.25
P0C0L4;P0C0L5	P0C0L4;P0C0L5	CO4A_HUMAN;CO4B_HUMAN	C4A;C4B_2	4478.57	3497.83	5165.04			594.613	258.368		
P0C0S5;Q71UI9	P0C0S5;Q71UI9	H2AV_HUMAN;H2AZ_HUMAN	H2AZ1;H2AZ2	75981.8	71236.7	37212.4	18634.4	37713.8	17752.6	5232.44	5758.99	3865.13
P0C0S8;P20671;Q16777;Q6FI13;Q	P0C0S8;P04908;Q6FI13;P20671;Q9BTM1;Q7L7L0;Q9	H2A1D_HUMAN;H2A1H_HUMAN;H2A1J_HU	H2AC12;H2AC14;H2AC17;H	209204	229175	137682	172797	227009	148473	61435.4	98377.5	56963.1
P0C221	P0C221	CC175_HUMAN	CCDC175		12871.7		3944.83					
P0C7P0	P0C7P0	CISD3_HUMAN	CISD3	6607.61	4655.43	3802.99	2838.71	1412.23	622.06	1848.4	2838.17	1428.79
P0DMV8;P0DMV9	P0DMV9;P0DMV8	HS71A_HUMAN;HS71B_HUMAN	HSPA1A;HSPA1B	30864.3	39339.1	29112.8	20411.3	16830.2	22536.9	23943.3	19685	25237
P0DP23;P0DP24;P0DP25	P0DP24;P0DP25;P0DP23	CALM1_HUMAN;CALM2_HUMAN;CALM3_H	CALM1;CALM2;CALM3	10595.5	9942.65	12027.3	16322.6	8646.66	8883.79	39224.5	17587.7	32476
P0DPB6	P0DPB6	RPAC2_HUMAN	POLR1D				3643.34				6641.04	
P0DPH7;P0DPH8;Q6PEY2	P0DPH7;Q6PEY2;Q71U36;P0DPH8	TBA3C_HUMAN;TBA3D_HUMAN;TBA3E_HU	TUBA3C;TUBA3D;TUBA3E	2037.13	4171.92	2653.75			1818.3	2292.89	3464.48	3101.63
P0DTE7;P0DTE8;P0DUB6	P0DTE7;P0DUB6;P0DTE8	AMY1A_HUMAN;AMY1B_HUMAN;AMY1C_	AMY1A;AMY1B;AMY1C	12151.9	3013.75	3592.04	2179.14	842.925	914.719	1939.27	3583.17	2296.81
P10109	P10109	ADX_HUMAN	FDX1						1156.13			
P10114	P10114	RAP2A_HUMAN	RAP2A							2677.96		2533.04
P10155	P10155	RO60_HUMAN	RO60	16816.5	14870	17990.5	18332.1	9122.71	10986.9	14442.7	12548.1	17349.3
P10301	P10301	RRAS_HUMAN	RRAS				4120.29	8846.47	9119.42	8782.08		10273.2
P10321	P10321	HLAC_HUMAN	HLA-C				4759.67	3248.58	3659.19	8801.17	12499.7	9801.27
P10398	P10398	ARAF_HUMAN	ARAF	6208.91	4603.69	9888.37	5237.59	4633.35	4554.38	9646.96	9292.18	11023.4
P10515	P10515	ODP2_HUMAN	DLAT	11054.7	23251.3	23887.6	109259	13826.4	129583	5944.5	6363.47	17303.9
P10589;P24468	P10589;P24468	COT1_HUMAN;COT2_HUMAN	NR2F1;NR2F2							661.289	1287.83	
P10599	P10599	THIO_HUMAN	TXN	28211.7	13074.4	18658.3	16480.7	15872.9	18672.8	18775.8	29150.1	15423.9

P10606	P10606	COX5B_HUMAN	COX5B	5981.26	4414.03	3940.98	3645.85	3921.09	2991.24	8364.69	4655.5	6707.73
P10619	P10619	PPGB_HUMAN	CTSA	14660.6	12002.4	15208.9	2639.2	2052.74	3306.89	3074.01	3524.2	2755.07
P10620	P10620	MGST1_HUMAN	MGST1	30246.5	37914.6	10522.4	5636.99			7929.98	4885.5	4472.72
P10644	P10644	KAP0_HUMAN	PRKAR1A	9760.26	7253.14	5329.82	6274.02	9084.37	10036	4975.71	4120.59	4070.77
P10768	P10768	ESTD_HUMAN	ESD				427.727	1194.33	5240.63	968.304		511.114
P10809	P10809	CH60_HUMAN	HSPD1	41930	19469.7	41445.2	38434.5	60500.8	76918	93134.8	54715.9	89176.9
P11021	P11021	BIP_HUMAN	HSPA5	106636	90909.3	78737.6	58153.4	53391.4	73334.5	41229.7	27448.7	39403.6
P11047	P11047	LAMC1_HUMAN	LAMC1	974.499			2300.9	878.835	1601.4	443.282		494.865
P11142	P11142	HSP7C_HUMAN	HSPA8	156316	107906	129885	186851	166579	228143	179685	141626	184948
P11166	P11166	GTR1_HUMAN	SLC2A1	7596.57	9325.57	9012.05	13163.8	61458.6	35778.4	130824	33225	124843
P11169	P11169	GTR3_HUMAN	SLC2A3		1051.08			7308.73	5503.88	10974.9	4951.9	8732.2
P11171	P11171	EPB41_HUMAN	EPB41						1581.61	3670.11		3643.13
P11172	P11172	UMPS_HUMAN	UMPS	14162.6	10749.2	13210.6	19778.7	11951.4	12356.4	16869.7	26682.2	18469.3
P11177	P11177	ODPB_HUMAN	PDHB	2878.04	5109.04	3945.96	9363.83	9080.86	16492.6	8813.94	4628.33	5560.68
P11182	P11182	ODB2_HUMAN	DBT	649947	678064	711581	847589	234182	225402	396252	896432	431924
P11216	P11217;P11216	PYGB_HUMAN	PYGB	8277.23	9104.81	10772.9	13409.7	15762.7	17341.1	13496.7	7523.3	12554.5
P11233	P11233	RALA_HUMAN	RALA			5231.74	6876.61	14996.5	10959	33534.7	7046.8	30846.1
P11234	P11234	RALB_HUMAN	RALB				2298.69	5835.74	5862.56	10785.1	2755.49	9204.1
P11274;Q12979	P11274;Q12979	ABR_HUMAN;BCR_HUMAN	ABR;BCR			1809.18	2907.08	1821.18	3629	4441.15	1727.09	5047.27
P11279	P11279	LAMP1_HUMAN	LAMP1	9652.49	11247.3	12406.9		6520.8	6511.41	10352.7	3466.75	11758.3
P11310	P11310	ACADM_HUMAN	ACADM	10959.8	8869.14	9937.42	11499.2	12801.9	8133.95	14886.7	21630.4	14358.1
P11387	P11387	TOP1_HUMAN	TOP 1.00	9300.89	8237.48	6828.78	6260	7010.05	4323.38	1699.04	4219.86	4208.87
P11388	P11388	TOP2A_HUMAN	TOP2A	2238.54	2407.53		2481.12	4261.19	1881	2184.56	3675.84	2085.8
P11413	P11413	G6PD_HUMAN	G6PD	16961.9	11395.5	16270.5	9176.06	9623.3	11470	10711.7	6864.94	11045
P11441	P11441	UBL4A_HUMAN	UBL4A	2327.78		2805.33	4244.42	2193.9	4688.59	2938.39	2283.97	3327.87
P11498	P11498	PYC_HUMAN	PC	878776	677136	936922	837446	200687	193065	869626	2.25502E+06	1.02646E+06
P11586	P11586	C1TC_HUMAN	MTHFD1	16139.1	12497.9	16740.3	19119.6	19287.5	19687.1	22656.8	21967.3	22747.4
P11717	P11717	MPRI_HUMAN	IGF2R	3477.07	4427.96	4889.73	5812.23	4276.06	4991.91	1917.23	761.931	1915.44
P11766	P11766	ADHX_HUMAN	ADH5					1334.89	2717.95	1867.58		2467.42
P11802	P11802	CDK4_HUMAN	CDK4	17248.5	22879.8	16841.1	6908.35	6969.94	4313.57	9319.78	6345.04	7307.9
P11908	P11908	PRPS2_HUMAN	PRPS2				3143.35	4245.02	5022.51	9122.92	7686.6	10867.8
P11940	P11940;Q4VXU2	PABP1_HUMAN	PABPC1	127197	66819.7	31535.9	69952.2	139785	41138.3	50266.4	54320.7	45071.4
P12004	P12004	PCNA_HUMAN	PCNA			4797.39	4954.96		4103.6	4414.78	5434.26	4071.35
P12035	P12035	K2C3_HUMAN	KRT3			29306.8	29393.1					
P12081	P12081;P49590	HARS1_HUMAN	HARS1	4814.18	2298.39	5992.15	5896.19	6831.98	12988.7	3857.14	3213.77	6810.39
P12235	P12235	ADT1_HUMAN	SLC25A4						3049.41	2749.37		7602.74
P12259	P12259	FA5_HUMAN	F5							31761.4	9998.72	32997.4
P12268	P12268	IMDH2_HUMAN	IMPDH2	80248.2	233087	15314	159566	339673	30320.4	26554	26620.2	23957.7
P12270	P12270	TPR_HUMAN	TPR				1453.23	2607.55	2022.11	3432.78	1368	3386.05
P12273	P12273	PIP_HUMAN	PIP	30863.2	21090.9	44742.9	16877.8	5838.39	11304.6	5639.04	20774.4	13161.2
P12429	P12429	ANXA3_HUMAN	ANXA3	2847.76		3601.93	3240.13	3190.83	6442.71	3796.92	2301.92	3003.84
P12532	P12532	KCRU_HUMAN	CKMT1B							1979.63	1322.01	1886.98
P12694	P12694	ODBA_HUMAN	BCKDHA	14827.6	14339.3	15111.2	7810.49	6493.4	6510.54	13153.8	17792.3	16411.2
P12814	P12814	ACTN1_HUMAN	ACTN1	15272.5	28797	16241.5	38504.2	33421.6	33422.6	17456.2	10082.1	15914.8
P12830	P12830	CADH1_HUMAN	CDH1							23423.7	5125.81	24370.5
P12882	Q9UKX2;P12882	MYH1_HUMAN	MYH1			624280				890616	151882	726646
P12883	P12883	MYH7_HUMAN	MYH7		5219.38	3311.19	11586.1	6968.93			6924.32	1540.22
P12931	P12931	SRC_HUMAN	SRC					4888.18	2128.85	4275.84	1442.86	3105.02
P12955	P12955	PEPD_HUMAN	PEPD	3047.04	1580.88	2670.79	1266.38	3427.58	8367.06	3124.95	2305.75	3874.36
P12956	P12956	XRCC6_HUMAN	XRCC6	#########	#########	71058.6	1.16938E+06	963237	54267	59244.1	738105	37069.6
P12980	P12980	LYL1_HUMAN	LYL1									1642.48
P13010	P13010	XRCC5_HUMAN	XRCC5	#########	#########	59401.5	2.58516E+06	1.30128E+06	68779.2	86000.9	870735	38170.5
P13073	P13073	COX41_HUMAN	COX4I1				5291.92			3260.81	10962.3	2954.05
P13473	P13473	LAMP2_HUMAN	LAMP2	11839.8	7002.57	18762.8	27394.6	11449.8	68214.3	15863	17494	11655.6
P13489	P13489	RINI_HUMAN	RNH1	14343.2		16197	11202.4	15646.8	10938.3	12696.5	7896.93	6991.28
P13533	P13533	MYH6_HUMAN	MYH6									
P13639	P13639	EF2_HUMAN	EEF2	30585.2	16016.7	38516.4	36833.5	41461.2	72337.7	31265.1	20174.1	29762.8
P13645	P13645	K1C10_HUMAN	KRT10	#########	1.2903E+06	2.54851E+06	929853	288513	615105	666364	1.07599E+06	1.17443E+06
P13646	P13646	K1C13_HUMAN	KRT13	19746.6		63235.9	55645.7		15801.4	26582.7	159689	41639.3
P13647	P13647	K2C5_HUMAN	KRT5	243128	213336	421874	177914	65995.4	117958	130626	289311	254902
P13667	P13667	PDIA4_HUMAN	PDIA4	10455		10918.9	5542.05	5648.81	9774.61	8915.3	5953.13	7737.52
P13674	P13674	P4HA1_HUMAN	P4HA1		4927	4622.75	7244.87	10316.5	7432.9	6009.32	7672.76	6380.85
P13688	P13688	CEAM1_HUMAN	CEACAM1				394.126	1098.52	979.323	6181.28	627.218	4756.24
P13693	Q56UQ5;P13693	TCTP_HUMAN	TPT1	3574.35		5683.8	8262.6	7594.8	26085	3736.33	2289.76	3939.68
P13716	P13716	HEM2_HUMAN	ALAD									
P13796	P13796	PLSL_HUMAN	LCP1			6349.2	25865.3	10536.7	31129.5	11970.7	10328.6	15466.8
P13797	P13797	PLST_HUMAN	PLS3	10270.5	8359.49	12350.9	12417.2	11519.6	20312	10116.6	5139.23	11157.2
P13798	P13798	ACPH_HUMAN	APEH		1188.36	3474.38	2565.28	3484.91	5640.55	3619.2	3649.23	2545.6
P13804	P13804	ETFA_HUMAN	ETFA	21383	28765.2	28346.2	46459.6	37995.1	20752.6	41180.7	73812.9	33301.4
P13805	P13805	TNNT1_HUMAN	TNNT1									
P13807	P13807	GYS1_HUMAN	GYS1			2866.77	2198.32	3607.68	4040.46	1556.09		1714.64
P13861	P13861	KAP2_HUMAN	PRKAR2A				3810.19	2238.89	916.844	1271.16	3306.35	1792.84
P13929	P13929	ENOB_HUMAN	ENO3	139768	16907.7	53883.4	33270.2	24792.7	68655.7	66422.2	29662.8	40281.2
P13984	P13984	T2FB_HUMAN	GTF2F2	3165.13		3065.26		3614.38	3869.13	2580.66		
P13987	P13987	CD59_HUMAN	CD59		5332		15892.2	10812.6	17833.1	11637.9	8103.38	11509.4
P13995	P13995	MTDC_HUMAN	MTHFD2					1876.05	4878.17	1888.44		1704.59
P14174	P14174	MIF_HUMAN	MIF	71669.8	8601.04	72182	32067	12951.1	52692.6	39950.1	23879.5	37806.1
P14222	P14222	PERF_HUMAN	PRF1							5536.87	4082.71	9153.35
P14314	P14314	GLU2B_HUMAN	PRKCSH	9475.1	5569.56	8352.7	11505.6	8583.65	16873.8	12183.5	6735.69	10537.8
P14324	P14324	FPPS_HUMAN	FDPS	2665.44			4211.04		3012.72	2481.64	1807.24	1892.91
P14373	P14373	TRI27_HUMAN	TRIM27	4307	3974.92	4110.77	3102.89	15451.3	10229	22037.7	1926.28	27576.6
P14550	P14550	AK1A1_HUMAN	AKR1A1	8515.65	4324.68	8244.49	4652.03	7897.43	23338.7	2990.51	2084.05	2471.86
P14618	P14618	KPYM_HUMAN	PKM	59052.8	34428.9	71617.9	38734.9	44805.7	102659	58694.5	30082.1	54764.4
P14625	P14625	ENPL_HUMAN	HSP90B1	22574.4	13685.1	25051.8	19879.3	19385	28281.5	22107.7	13038.9	24809.7
P14635	P14635	CCNB1_HUMAN	CCNB1				910.593	6508.11	4581.41	2924.82	1593.52	1813.56
P14649	P14649	MYL6B_HUMAN	MYL6B	1009.17			5067.23	8093.32	4657.53	2916.36	3219.34	2088
P14678;P63162	P14678;P63162	RSMB_HUMAN;RSMN_HUMAN	SNRPB;SNRPN	5511.75	5997.43	4537.86	8556.9	26703.4	4558.3	13911.5	13379.2	10994.5
P14735	P14735	IDE_HUMAN	IDE	6702.63		6928.77	5788.12	7831.95	6127.72	7825	5796.06	6853.53
P14866	P14866	HNRPL_HUMAN	HNRNPL	4086.7		2818.09	5114.41	4238.57	2015.72	1625.51	2768.24	1579.54
P14868	P14868	SYDC_HUMAN	DARS1	13883.5	13301.9	11743.4	17802.7	15684.6	15734.5	15302	14652.4	15711
P14923	P14923	PLAK_HUMAN	JUP	44022.4	25699.2	52680.1	42969.4	39111.3	34457.5	49110.8	28189.9	48778.2
P15104	P15104	GLNA_HUMAN	GLUL							2490.84	1011.53	3222.36
P15121	P15121	ALDR_HUMAN	AKR1B1		3141.22	9260.22		3274.98	12933.4	2945.3	1826.54	
P15144	P15144	AMPN_HUMAN	ANPEP			2410.66	8981.56	9130.85	9563.84			
P15151	P15151	PVR_HUMAN	PVR	1037.22	721.53	959.482	938.239	3997.88	4213.58	6299.78	616.605	5072.08
P15153	P15153	RAC2_HUMAN	RAC2				8949.07	2532.43	10518.1	2612.96	2443.71	3888.41
P15170	P15170	ERF3A_HUMAN	GSPT1	3833.72	5664.08	6277.32	6521.29	10569.6	18797.9	5941.83	2792.23	6409.37
P15259	P15259	PGAM2_HUMAN	PGAM2						3173.61			
P15311	P15311	EZRI_HUMAN	EZR	7977.8		7940.05	14161.9	16069.3	26902.4	15633.5	5885.84	16581.8
P15328	P15328	FOLR1_HUMAN	FOLR1				3988.99	4199.52	3281.18	10065	3815.76	10569.8
P15428	P15428	PGDH_HUMAN	HPGD									
P15531	P15531;O60361;P22392	NDKA_HUMAN	NME1	59830.9	27156.5	70785	95274.5	66436.9	164634	59791.9	36214.5	65493.9
P15559	P15559	NQO1_HUMAN	NQO1	37280.9	26643.2	37389.9	8555.57	2789.99	5997.09	27602.8	16380.2	23638.5
P15586	P15586	GNS_HUMAN	GNS	3724.16		4243.54	3324.88		2056.39	5838.07		
P15880	P15880	RS2_HUMAN	RPS2	53424.2	63406.3	50904.9	38783.9	78044.2	57476.2	53888.9	31317.4	44499.3
P15882	P15882	CHIN_HUMAN	CHN1				1116.25	3215.21	2383.38	956.531		
P15924	P15924	DESP_HUMAN	DSP	53308.1	35194	64957.2	33248.5	19603.9	17122.4	78713.5	42105.5	66004.6
P15927	P15927	RFA2_HUMAN	RPA2	35134.2	270012	7690.69	17096.2	21042	6534.95	7244.27	32237.8	
P16070	P16070	CD44_HUMAN	CD44	14484.8	12313.4	9545.94	10283.5	20992.4	36936.1	5992.02	3203.61	4496.09
P16144	P16144	ITB4_HUMAN	ITGB4		7707.62	7820.25		4581.24		59085.3	21066	44451.9
P16152	P16152	CBR1_HUMAN	CBR1	4006.38	1999.68	3478.83	2125.36	731.656	1432.58	790.838		
P16219	P16219	ACADS_HUMAN	ACADS						587.55			
P16220	P16220	CREB1_HUMAN	CREB1									
P16278	P16278	BGAL_HUMAN	GLB1	45296.7	48301.3	51196.6	1316.26	7332.89	8099.68	6208.68	2048.22	5264.44
P16401	P16401	H15_HUMAN	H1-5	14699.7	12986.8	8827	3176.05	6526.61	3923.23	3271.92	2774.61	2225.05
P16403	P16402;P10412;P16403	H12_HUMAN	H1-2	37272.2	36915.1	21431.9	19200.8	17928.9	9869.72	14828.5	15650.9	9484.85
P16422	P16422	EPCAM_HUMAN	EPCAM							10901.7	6543.87	9835.9
P16435	P16435	NCPR_HUMAN	POR						435.254	1161.72		1770.03
P16444	P16444	DPEP1_HUMAN	DPEP1							24651.1	8775.08	18458.2
P16615	Q93084;P16615	AT2A2_HUMAN	ATP2A2	9002.81	7797.77	11675.5	10775.6	7126.2	5950.71	11530.5	11784.9	12749.9
P16671	P16671	CD36_HUMAN	CD36				6253.56				5094.12	
P16930	P16930	FAAA_HUMAN	FAH			2179.32	1360.58	3387.18	8703.69	1621.85		988.45
P16949	P16949	STMN1_HUMAN	STMN1	12149	4309.68	13158.9	12720.9	9919.77	11986.7	13541	10831.6	16182.5
P16989	P16989	YBOX3_HUMAN	YBX3	4762.3	6756.51	1505.28	5801.88	4979.2	3822.43	3318.26	7910.33	2208.56
P17066	P48741;P17066	HSP76_HUMAN	HSPA6				15260.6				7731.2	3684.99
P17096	P17096	HMGA1_HUMAN	HMGA1	60226.2	20800.8	19757.7	71381.4	159046	74815.6	18870.8	31487.4	24995.5
P17174	P17174	AATC_HUMAN	GOT1	4853.55		4796.29	2329.73	4765.33	7137.2	2044.24		2146.5
P17252	P17252	KPCA_HUMAN	PRKCA				4151.83	5968.33	4353.02	5566.65	1852.23	5532
P17301	P17301	ITA2_HUMAN	ITGA2	1604.49				3287.64	3231.92	1778.58		1103.99
P17480	P17480	UBF1_HUMAN	UBTF	7562.48	23768.1	7885.94	3552.32	12990.2	3476.59	3578.66	6088.71	4333.83
P17535	P17535	JUND_HUMAN	JUND									
P17540	P17540	KCRS_HUMAN	CKMT2									
P17544	P17544	ATF7_HUMAN	ATF7					3687.55	1987.77			
P17600	P17600	SYN1_HUMAN	SYN1				9824.35		1715.56	7554.92	4610.29	12104.5
P17612	P22694;P17612	KAPCA_HUMAN	PRKACA	6782.65	7828.47	2939.89	1634.15	4048.59	4098.37	3978.21	1231.37	4550.09
P17655	P17655	CAN2_HUMAN	CAPN2	11862.6	6620.18	5944.9	10678.2	6016.29	6125.75	6262.57	10825.8	8593.09
P17661	P17661	DESM_HUMAN	DES	26484.3	29683.7	13180.3			15002.1	29182.1		
P17676	P17676	CEBPB_HUMAN	CEBPB	4162.39			2248.39	2295.9				
P17677	P17677	NEUM_HUMAN	GAP43							4554.2	979.014	4757.68
P17812	P17812;Q9NRF8	PYRG1_HUMAN	CTPS1	15358.5	16198.1	14330	30656.8	33544	25735	23146.9	21938.6	25269.6
P17813	P17813	EGLN_HUMAN	ENG	1110.6		2482.97	2089.61	5142.84	5347.44			
P17844	P17844	DDX5_HUMAN	DDX5	50844.4	53322.1	53079.1	41779.6	66789.4	33581	58759	45531.3	56099.5
P17858	P08237;P17858	PFKAL_HUMAN	PFKL	18030.5	19663.3	17716.9	10436.4	7524.72	8483.66	13889	11935.4	14188
P17900	P17900	SAP3_HUMAN	GM2A			4122.75	1970.05				2602.18	
P17931	P17931	LEG3_HUMAN	LGALS3	13315.9	7793.24	13460.5	5433.24	9621.52	14479.4	338398	30821.3	248200
P17980	P17980	PRS6A_HUMAN	PSMC3	12243.1	8173.76	13233.2	15460.7	17755	14241.8	10452.7	8441.17	11776.8
P17987	P17987	TCPA_HUMAN	TCP1	33684	18575.3	33348.9	47627.5	44625.2	44679	53306.3	58095.3	52878.2
P18031	P18031	PTN1_HUMAN	PTPN1				2885.01	4450.14	3081.88	3999.53	1778.89	4317.59
P18074	P18074	ERCC2_HUMAN	ERCC2					3848.34	1379.85			
P18077	P18077	RL35A_HUMAN	RPL35A	31284.5	44147.9	18859.4	31786.3	93203.9	40778.1	37370.6	33428	40298.5
P18085	P18085	ARF4_HUMAN	ARF4	9990.72	9036	14454.8	31273.7	15464.9	7112.24	4287.75	14489.3	3588.25
P18124	P18124	RL7_HUMAN	RPL7	37696.4	57103	45161.2	58388.4	70204	51164.3	56958.6	47606.2	51317.3
P18206	P18206	VINC_HUMAN	VCL	10411.1	6682.09	9897.85	10634.1	12972	22102.9	11655.9	4751.26	11553
P18615	P18615	NELFE_HUMAN	NELFE					1222.05	1704.9			
P18621	P18621	RL17_HUMAN	RPL17	5133.9	12071.6	6396.26	6504.46	23513.3	17939.9	6920.41	10146.9	5555.19
P18669	P18669	PGAM1_HUMAN	PGAM1	10882.6	3825.39	14264	5782.14	15653.8	40716.3	15553.2	3974.83	18632.3
P18754	P18754	RCC1_HUMAN	RCC1		1973.91	1928.03	1156.34	2619.38	3811.29	2060.3		1986.94
P18858	P18858	DNLI1_HUMAN	LIG1				1349.45	1662.45	2014.57	1496.79		1458.39
P18887	P18887	XRCC1_HUMAN	XRCC1	23577.4	44539.1		29809.6	34590.5	4891.12	4003.28	5630.28	2654.28
P19012	P19012	K1C15_HUMAN	KRT15			29540.5			3491.35	6972.71	79604.8	6132.75
P19013	P19013;Q86Y46;Q3SY84	K2C4_HUMAN	KRT4	2200.71	730.45	16307.8	5750.94		4059.71	6378.1	49555.8	5571.42
P19022	P19022	CADH2_HUMAN	CDH2				3861.56	13949.4	16294.4			
P19086	P19086	GNAZ_HUMAN	GNAZ						10186.5	20154.7		22690.1
P19174	P19174	PLCG1_HUMAN	PLCG1					4930.72	5691.51	3460.81		1655.37
P19237	P19237	TNNI1_HUMAN	TNNI1									
P19256	P19256	LFA3_HUMAN	CD58					3711.07	2580.02	6641.07	1889.2	5374.13
P19338	P19338	NUCL_HUMAN	NCL	18589.2	14044.4	15846.7	18308.7	19785.5	19570.7	33881.3	17494.5	30080
P19367	P19367	HXK1_HUMAN	HK1	2937.02		5289.45	4860.3	7305.72	9252.59	2821.95		2599.23
P19388	P19388	RPAB1_HUMAN	POLR2E									
P19404	P19404	NDUV2_HUMAN	NDUFV2	3354.63	3500.12	4708.49	3448.29	3884.18	6716.76	10129.7	8149.27	8136.04
P19525	P19525	E2AK2_HUMAN	EIF2AK2	5397.55	3598.23	5336.12	3249.64	2449.86	1874.04	2696.78	2610.52	3099.08
P19532	P19532	TFE3_HUMAN	TFE3	1991.1			2048.17					
P19623	P19623	SPEE_HUMAN	SRM	15359.8	12267.2	17668.4	19718.5		10285.8	16183.1	8983.69	17417.2
P19634	P19634	SL9A1_HUMAN	SLC9A1	670.804	786.876	1040.49	296.742	539.095	284.526	1693.56		1051.41
P19838	P19838	NFKB1_HUMAN	NFKB1					1168.53	723.098			
P19957	P19957	ELAF_HUMAN	PI3			1036.23	627.769				299.464	
P20020	P23634;P20020	AT2B1_HUMAN	ATP2B1	6252.69	5880.18	6259.15	4774.91	3105.7	2859.66	2254.61	2276.67	2262.13
P20042	P20042	IF2B_HUMAN	EIF2S2	4406.68			4422.02	6501.15	6793.06	4683.28	3903.76	3879.35
P20073	P20073	ANXA7_HUMAN	ANXA7	9290.13	3124.49	8306.82	5310.42	4212.62	5889.6	7504.98	5579.88	5874.7
P20226;Q6SJ96	Q6SJ96;P20226	TBPL2_HUMAN;TBP_HUMAN	TBP;TBPL2				1416.31					
P20290	P20290	BTF3_HUMAN	BTF3		940.669	1153.65	654.341	2870.15	4137.28	1695.04	581.723	1979.2
P20339	P20339	RAB5A_HUMAN	RAB5A				3049.65	2316.21	4876.69	4714.38	2656.65	5076.24
P20340	P20340	RAB6A_HUMAN	RAB6A	5027.38		6680.31	6026.63	29271.5	3431.75	10771.8	12640.1	8011.41
P20585	P20585	MSH3_HUMAN	MSH3	22549.3	41129.5	2574.45	14581.1	11443.7	2432.98	3362.59	11647.4	3203.92
P20591	P20591	MX1_HUMAN	MX1									
P20592	P20592	MX2_HUMAN	MX2									
P20618	P20618	PSB1_HUMAN	PSMB1	5637.08	4132.87	7338.94	3980.93	3150.86	6937.69	5304.18	3538.18	5941.07
P20700	P20700	LMNB1_HUMAN	LMNB1	6324.55	5263.5	5862.64		1699.32	2149.32	2002.25		
P20810	P20810	ICAL_HUMAN	CAST	3724.99		2556.12	4669.97	5681.62	6440.42	2402.36	3779.03	3025.63
P20839	P20839	IMDH1_HUMAN	IMPDH1	4958.32	20506.9	1563.36	5172.47	18229	3132.74	3242.74	1776.98	1194.8
P20908	P20908	CO5A1_HUMAN	COL5A1					3323.78				
P20930	P20930	FILA_HUMAN	FLG	21430.4	18688.3	24441.3	15156.4	15136.2	14088.2	19044.4	12696.3	21024.9
P20936	P20936	RASA1_HUMAN	RASA1					1698.7	2204.1	1275.89		
P21127;Q9UQ88	Q9UQ88;P21127	CD11A_HUMAN;CD11B_HUMAN	CDK11A;CDK11B	4903.03	13557.3	3262.94	6644.22	8187.39	3668.73	3928.75	7102.78	6385.13
P21281	P21281	VATB2_HUMAN	ATP6V1B2	6344.72	3573.69	8191.14	5378.69	7719.15	11609.6	7030.12	4547.88	7079.34
P21291	P21291	CSRP1_HUMAN	CSRP1	9392.05		10960	4012.6	3481.08	14393.8	2590.29		2821.96
P21333	P21333	FLNA_HUMAN	FLNA	43833	32637.5	47877.2	47690.9	37198.1	47602.7	32156.4	31491.4	26254.4
P21359	P21359	NF1_HUMAN	NF1		3210	1077.69	1320.68	2649.9	2387.78	1040.5	802.588	816.489
P21399	P21399	ACOHC_HUMAN	ACO1	7025.1	3911.73	13302.4	3348.1	3391.88	5650.28	2935.47		3153.26
P21579	P21579	SYT1_HUMAN	SYT1							7991.49	8568.54	7194.81
P21589	P21589	5NTD_HUMAN	NT5E	16212.7	16495.4	15321.7	37940.6	31571.8	42718.9	6065.2	5491.19	4980.49
P21695	P21695	GPDA_HUMAN	GPD1					471.322				
P21796	P21796	VDAC1_HUMAN	VDAC1	19231.5	25222.9	19156	3134.01	7292.6	6968.66	11175.2	6019.56	10494.4
P21912	P21912	SDHB_HUMAN	SDHB				2803.77			5627.91	4612.44	7642.69
P21926	P21926	CD9_HUMAN	CD9					4489.07	1309.9	6128.6		6197
P21953	P21953	ODBB_HUMAN	BCKDHB	6990.53		10621.3	8833		3088.26	9892.29	13104.2	13498.2
P21964	P21964	COMT_HUMAN	COMT			2745.12		2135.84	3538.48	1675.09		1579.98
P21980	P21980	TGM2_HUMAN	TGM2	12657.6	11775.3	10430.6	37112.2	31243.5	38419.3	1406.91		886.541
P22059	P22059	OSBP1_HUMAN	OSBP					1111.13	1314.5	824.123		935.048
P22061	P22061	PIMT_HUMAN	PCMT1				1792.58	5007.18	6474.88	2840.9	865.711	3062.86
P22087	P22087	FBRL_HUMAN	FBL	6566.22	11157.3	3986.28	2133.37	8165.21	2821.4	6736.66	5043.12	6521.37
P22090	P22090;Q8TD47	RS4Y1_HUMAN	RPS4Y1	10345.8	9601.28	9158.24						
P22102	P22102	PUR2_HUMAN	GART	6361.13	3769.77	7155.41	15588.8	19409.7	31177.3	16783.2	16686.9	15046.5
P22234	P22234	PUR6_HUMAN	PAICS	6559.57	4357.37	8729.81	10161.8	10972.9	19523.8	29863.2	10255.4	19803.8
P22307	P22307	SCP2_HUMAN	SCP2	4739.14			2851.87		4085.62	3652.89		
P22314	P22314	UBA1_HUMAN	UBA1	10071.5	9316.19	10244.8	10914.2	9941.49	15443	16360.4	10290.9	12926.9
P22392	O60361;P22392	NDKB_HUMAN	NME2	9819.74	5058.7	12245.8	13149.6	5333.56	20147.1	10562.6	8773.4	11083.2
P22455	P22455	FGFR4_HUMAN	FGFR4		319094		14213.6	59145.1	58357.6	153475	10865.8	165967
P22528	P22528;P35321	SPR1B_HUMAN	SPRR1B	10932.2	14916.9	16137.3	20508		3136.1	8086.84	22495.6	11508.4
P22532;P35325	P22532;P35325	SPR2B_HUMAN;SPR2D_HUMAN	SPRR2B;SPRR2D			4379.92	7472.13		4817.23	2303.83	7268.96	5050.73
P22570	P22570	ADRO_HUMAN	FDXR				752.191	700.456	919.054	458.848	416.481	392.696
P22626	P22626	ROA2_HUMAN	HNRNPA2B1	9730.7	5920.55	11353.2	16166.9	96777.1	35077.7	15822.1	10777.4	14820.8
P22694	P22694	KAPCB_HUMAN	PRKACB	944.144	5928.87	733.077	386.495	9401.04	4488.72	2101.15	535.392	3237.71
P22695	P22695	QCR2_HUMAN	UQCRC2	3663.46	2893.17	3256.1	3288.74	4789.65	4587.99	9555.55	8164.76	9665.83
P22735	P22735	TGM1_HUMAN	TGM1	21661.6	13617.5	25313	21419.4	19994.6	6551.31	11762.1	15517.2	10684.4
P22830	P22830	HEMH_HUMAN	FECH	4690.15	6857.57			4644.1	3795.7	2295.78		2270.65
P23109	P23109	AMPD1_HUMAN	AMPD1									
P23193	Q15560;P23193	TCEA1_HUMAN	TCEA1				3586.27	3354.66	3418.91	1528.79		1636.61
P23229	P23229	ITA6_HUMAN	ITGA6				2646.89	3425.91	3409.98	16027.3	5401.09	11433.6
P23246	P23246	SFPQ_HUMAN	SFPQ	18328.4	14848.5	15191.7	21595	473317	22568.6	15181.8	15026.1	14354.3
P23258	P23258;Q9NRH3	TBG1_HUMAN	TUBG1	6058.92		7467.06	9592.31	4615.99	4236.34	5471.18	16102.9	5706.45
P23284	P23284	PPIB_HUMAN	PPIB	10748.9	3803.6	13399.2	6532.8	13367.3	22858.8	8486.44	4686	7025.03
P23381	P23381	SYWC_HUMAN	WARS1	12569.4	6151.57	9436.04	4532.46	8058.6	16749.1	5165.32	2361.37	5752.72
P23396	P23396	RS3_HUMAN	RPS3	82920.5	90624.3	73963.4	116080	131586	79042.5	91429.9	120895	84921.5
P23458	P23458	JAK1_HUMAN	JAK1	8822.73	7095.52	14851.3	2898.82	3214.92	3728.93	4174.71	4484.32	9538.38
P23470	P23470	PTPRG_HUMAN	PTPRG					7602.2	7473.83			
P23490	P23490	LORI_HUMAN	LORICRIN	6000.67	5765.98	7385.8	4054.32		1216.82	3083.79	4248.08	4579.76
P23526	P23526	SAHH_HUMAN	AHCY	16516.9	17717.5	15917.2	22367.8	15198.6	31580.1	46581.8	23102.4	48256.6
P23527;P33778;Q16778;Q8N257	P57053;Q99879;Q5QNW6;Q16778;P62807;O60814;P2	H2B1B_HUMAN;H2B1O_HUMAN;H2B2E_HU	H2BC17;H2BC21;H2BC26;H2	121262	94016.1	100538	155399	111123	69112.8	33622.8	90688.9	30193.8
P23528	Q9Y281;P23528	COF1_HUMAN	CFL1	22622.1	15896.6	28328	28500.4	52214.7	44294.5	17683.6	20192.3	16955.4
P23588	P23588	IF4B_HUMAN	EIF4B				4845.91	5197.9	10284.3	6528.19	5111.41	7996.37
P23743	P23743	DGKA_HUMAN	DGKA				49779.5					
P23786	P23786	CPT2_HUMAN	CPT2						1088.82	973.017		675.592
P23919	P23919	KTHY_HUMAN	DTYMK			4375.7	4045.19		3670	2584.36	2555.56	2826.77
P23921	P23921	RIR1_HUMAN	RRM1				4497.45	4033.37	6833.25	8119.41	5703.92	5382.77
P24158	P24158	PRTN3_HUMAN	PRTN3	5828.15		23888	4366.07		2344.48	3458.11	4434.79	3451.4
P24278	P24278	ZBT25_HUMAN	ZBTB25									
P24534	P24534	EF1B_HUMAN	EEF1B2	4829.18	3553.16	5241.6	5811.65	8564.21	5583.39	4396.09	5278.29	7546.62
P24539	P24539	AT5F1_HUMAN	ATP5PB	7124.49	6507.86	7198.27			1139.56	4061.58		5471.09
P24666	P24666	PPAC_HUMAN	ACP1						3287.01			
P24752	P24752	THIL_HUMAN	ACAT1	6973.62	7776.85	8001.58	6886.64	4583.15	7254.36	6754.31	6345.84	6173.95
P24844	P24844	MYL9_HUMAN	MYL9	8740.4	8023.85	4076.61	7193.3	62157.6	26458.3	7042.46	4127.58	
P24928	P24928	RPB1_HUMAN	POLR2A				2943.42	4704.77	2641.5	4925.58	3756.72	4518.04
P24941	P24941	CDK2_HUMAN	CDK2				960.029	4450.44	2832.45	2406.62	1380.86	1041.69
P25205	P25205	MCM3_HUMAN	MCM3	6411.81	8402.65	6301.12	5188.7	4530.28	4919.26	7805.33	5647.45	6730
P25311	P25311	ZA2G_HUMAN	AZGP1	14910.5	13170.8	21012.1	13158.8	23855.3	11224.6	7273.55	13927.5	11283.3
P25398	P25398	RS12_HUMAN	RPS12	110045	98132	64786	80097.9	84998.1	69125.1	96155.4	78494.4	75037.2
P25685	P25685	DNJB1_HUMAN	DNAJB1	5218.92		10658.7	2750.74	3512.97	4871.4	3076.67	3645.26	3570.63
P25705	P25705	ATPA_HUMAN	ATP5F1A	56803.1	45857.6	56079.9	26438.3	19162	15251.1	32274.4	34765.1	30068.5
P25786	P25786	PSA1_HUMAN	PSMA1	6516.61	4620.31	7498.33	5971.42	5012.51	10104.2	5396.75	4298.09	6476.07
P25787	P25787	PSA2_HUMAN	PSMA2	2953.57	2803.13	3596.13	2518.27	3827.71	6871.61	5198.09	3574.83	5325.96
P25788	P25788	PSA3_HUMAN	PSMA3	3567.4		6681.7	5946.04	9149.45	15685.4	6168.94	6321.06	8168.73
P25789	P25789	PSA4_HUMAN	PSMA4	6623.95	4809.16	10851	4247.38	4604.51	8368.92	10299.7	3306.63	8708.51
P26006	P26006	ITA3_HUMAN	ITGA3				4755.24	7535.87	7044.62			1035.32
P26038	P26038	MOES_HUMAN	MSN	50011.9	24763.6	54191.5	34852.7	38703.2	76845.7	48200.8	20713.2	49827.5
P26196	P26196	DDX6_HUMAN	DDX6	16327.3	12985.3	15484.7	12010.6	26630.3	27629.7	37801.2	13879.4	31760.8
P26232	P26232	CTNA2_HUMAN	CTNNA2					13763.8	7691.21	7831.89		3368.4
P26358	P26358	DNMT1_HUMAN	DNMT1					5013.36	3930.06	3022.17		1897.53
P26368	P26368	U2AF2_HUMAN	U2AF2	20029.1	24798.7	15241.9	14601.6	83497.4	14508.7	14272.1	16532.1	13453.8
P26373	P26373	RL13_HUMAN	RPL13	32836.4	34833	34360.3	41207.9	40588.1	36228.5	40547	29610.3	36535.1
P26440	P26440	IVD_HUMAN	IVD			1677.93	3308.5	1261.88	2291.41	3264.97	3514.08	2957.87
P26447	P26447	S10A4_HUMAN	S100A4				10052.7	7263.27	7859.06	14781.2	17265.8	16512.1
P26583	P26583	HMGB2_HUMAN	HMGB2	7655.78	9180.65	7663.32	5368.03	5782.73	7772.06	5412.62	4764.9	6428.01
P26599	P26599;O95758	PTBP1_HUMAN	PTBP1	23693.9	12017.8	13595.4	15240.6	34186.1	17308.6	25895.3	16008.6	26516.7
P26639	P26639;A2RTX5	SYTC_HUMAN	TARS1	12070.7	23987	10913.5	13581.2	6292	13321.4	8929.77	6658.55	14117.2
P26640	P26640	SYVC_HUMAN	VARS1	7873.82	3936.02	7826.45	11671	12676.5	13258.8	11256	7938.77	10768.6
P26641	P26641	EF1G_HUMAN	EEF1G	99127.8	68996.6	72672.3	124272	142482	180407	207293	144472	187789
P26885	P26885	FKBP2_HUMAN	FKBP2									
P27105	P27105	STOM_HUMAN	STOM	13323.1	11153.7	19118.5	2213.42	1581.11	1999.19	1090.91	1831.17	1111.74
P27144	P27144;A0A8I5KW96	KAD4_HUMAN	AK4	6570.85	4146.02	6353.72	10914	10925	13730.3	6803.94	5317.94	6918.47
P27348	P27348	1433T_HUMAN	YWHAQ	2013.76		5101.34	2448.47	3421.96	10488.6	5286.61	1067.82	6570.74
P27361	P27361;P31152;Q16659;P28482	MK03_HUMAN	MAPK3	5607.88	5738.75	6709.07	3401.81	4848.22	8510.41	4671.4	3489.18	5088.94
P27448	P27448	MARK3_HUMAN	MARK3				2101.1			4111.35	2132.79	3761.28
P27635	P27635;Q96L21	RL10_HUMAN	RPL10	31448.7	39025.1	34923.4	27545.4	60854.4	51233.6	37303.3	25167.3	41539.3
P27694	P27694	RFA1_HUMAN	RPA1	112338	536750	16590.8	55820.7	56745.3	17591.4	13505.7	114422	7695.42
P27695	P27695	APEX1_HUMAN	APEX1	3332.51	3705.32	3629.29	2267.83	3137.32	6412.54	3672.91	1899.29	3668.17
P27708	P27708	PYR1_HUMAN	CAD	10858.9	10691.4	7253.32	27698.5	20051.8	16755	31678.3	43066.6	28289
P27797	P27797	CALR_HUMAN	CALR	9825.34	3790.93	11174.3	5644.51	5259.7	8580.25	4984.91	4809.18	5983.37
P27816	P27816	MAP4_HUMAN	MAP4	16345.4	11415.3	12867.9	10644.4	8696.92	11007.9	5878.89	6556.6	6431.67
P27824	P27824	CALX_HUMAN	CANX	11900.5	6793.36	8558.36	5851.35	7854.74	8451.12	9393.4	4988.78	9909.05
P28062	P28062	PSB8_HUMAN	PSMB8			3286.03			2105.99			2439.66
P28066	P28066	PSA5_HUMAN	PSMA5	3449.27	3731.04	4668.8	3026.25	2808.08	5058.85	5748.52	3673.4	6699.9
P28070	P28070	PSB4_HUMAN	PSMB4	3219.05		5259.78			181.094			624.788
P28072	P28072	PSB6_HUMAN	PSMB6	6528.28		6760.76	5676.34	4789.24	9819.36	5687.35	5253.23	6128.15
P28074	P28074	PSB5_HUMAN	PSMB5	3838.77	3040.38	8757.48	3158.24	7078.55	17592.8	7915.07	2506	9191.18
P28288	P28288	ABCD3_HUMAN	ABCD3			1641.84	3129.92	3619.45	1026.5	1892.09	2443.89	2330.02
P28290	P28290	ITPI2_HUMAN	ITPRID2	6868.2	4755.19	5546.15	4686.39	3929.31	3475.42	10073.4	7475.09	9579.96
P28300	P28300	LYOX_HUMAN	LOX	1059.88		934.321						
P28325	P28325	CYTD_HUMAN	CST5			845.913						
P28330	P28330	ACADL_HUMAN	ACADL									
P28331	P28331	NDUS1_HUMAN	NDUFS1	2229.05		2857.75	1825.62	7479.89	5847.92	4500.24	4168.29	4092.78
P28340	P28340	DPOD1_HUMAN	POLD1			2543.78	11828.7	12104	9009.34	18892.3	18355.4	15808.3
P28347	Q99594;P28347;Q15561	TEAD1_HUMAN	TEAD1				982.185	3690.7	1084.76	810.221		1686.24
P28482	P28482	MK01_HUMAN	MAPK1					3157.98	5196.16	1600.65		1984.93
P28749	P28749	RBL1_HUMAN	RBL1									493.11
P28799	P28799	GRN_HUMAN	GRN	30926.7	34487.5	22177.2				3495.2		
P28838	P28838	AMPL_HUMAN	LAP3	7191.09		6547.08	4311.76	3850.43	4509.59	4091	4582.44	3212.99
P29144	P29144	TPP2_HUMAN	TPP2				2150.22	6153.72	3894.93	4371.52	3507.72	4619.37
P29218	P29218	IMPA1_HUMAN	IMPA1	1813.68		1933.12	2668.13	1382.39	3683.61	2500.93	3258.19	2595.34
P29317	P29317	EPHA2_HUMAN	EPHA2				4422.69	5851.02	6054.55	12158.7	5716.32	9260.34
P29323	P29323;P54762	EPHB2_HUMAN	EPHB2							5992.24		1690
P29372	P29372	3MG_HUMAN	MPG	4872.25	10255.2		6098.55	15975.9	4689.84	2753.39	8258.34	
P29373	P29373	RABP2_HUMAN	CRABP2			1224.04	1535.42	805.263	2928.85	698.68	964.452	779.015
P29401	P29401	TKT_HUMAN	TKT	28168.9	11664.6	25693	8632.74	11624.5	28500.5	14058.2	6255.42	15728.8
P29508	P29508	SPB3_HUMAN	SERPINB3	24908.8	21109.7	32520.3	20252.2	4822.8	13195.9	14197.9	21377.8	26756.6
P29558	P29558	RBMS1_HUMAN	RBMS1	62317.7	35704.8	5246.36	44293.7	12811.5	1885.98		29484.6	
P29692	P29692	EF1D_HUMAN	EEF1D	10304.5	8285.74	6459.28	16228.1	4026.38	8544.1	17931.6	8907.84	11370.2
P29728	P29728	OAS2_HUMAN	OAS2		3276.63	1713.06	1866.93	1149.09	1034.85			
P29966	P29966	MARCS_HUMAN	MARCKS	8981.75	3910.52	6291.89	13909.7	21948.2	24898.1	30252.1	13352.2	25985.4
P29992	P29992	GNA11_HUMAN	GNA11	2208.71		1912.77	12944.9	25449.9	25630.2	49861.7	11344.3	43218.9
P30038	P30038	AL4A1_HUMAN	ALDH4A1				2550.87			2392.13	4752.41	2088.71
P30040	P30040	ERP29_HUMAN	ERP29	12034.8	3496.32	9841.33	9367.64	9275.14	22632.4	4002.42		6358.88
P30041	P30041	PRDX6_HUMAN	PRDX6	22227.9	14809.7	32123.1	9762.96	11808.2	31489	13359.4	6124.99	14221.4
P30043	P30043	BLVRB_HUMAN	BLVRB	7446.49		6143.58			2977.7			
P30044	P30044	PRDX5_HUMAN	PRDX5	8122.86		6159.46	6666.08	3556.15	13696.2	2404.17		4406.94
P30046	A6NHG4;P30046	DOPD_HUMAN	DDT	4878.88		4893.96	2681.87	4167.14	10609.5	2849.65	2611.31	2051.15
P30048	P30048	PRDX3_HUMAN	PRDX3	19399.7	9529.79	23658.4	11635.3	7885.12	28473.5	25053.7	22375.3	20677.4
P30050	P30050	RL12_HUMAN	RPL12	27654.6	23300.3	22370.7	37595.6	20802	18870.5	22563.3	28680.9	23409.4
P30084	P30084	ECHM_HUMAN	ECHS1	6633.64	2002.59	9894.74	2639.17	3187.94	12276.3	8133.69	2308.67	9015.7
P30085	P30085	KCY_HUMAN	CMPK1			4423.76	3137.76	2635.95	3107.84	1915.75	2755.61	2977.45
P30086	P30086	PEBP1_HUMAN	PEBP1	3273.09		7890.11	3336.75		2779.18	2109.09	3093.08	2928.61
P30101	P30101	PDIA3_HUMAN	PDIA3	26211.7	7351.11	25411.6	16016.8	14927.2	34088.1	21891.8	9309.88	20377.3
P30153	P30153	2AAA_HUMAN	PPP2R1A	7451.43	8226.9	8545.85	11053.2	8032.19	14979	11726.9	9012.16	10878.2
P30154	P30154	2AAB_HUMAN	PPP2R1B	1273.28		839.977	3011.98	4568.18	5463.05	4515.9	1552.62	1662.76
P30405	P30405	PPIF_HUMAN	PPIF				4966.54		7372.4	3259.78		2975.56
P30419	P30419	NMT1_HUMAN	NMT1			986.972		2445.17	3002.71	1324.24	1337.61	1449.01
P30519	P30519	HMOX2_HUMAN	HMOX2					2312.18	2944.41	3033.3		1358.78

P30520	P30520	PURA2_HUMAN	ADSS2				4252.14		1726.81	3299.38		2527.67
P30530	P30530	UFO_HUMAN	AXL		1405.59	1801.01		1447.05	2145.5			
P30566	P30566	PUR8_HUMAN	ADSL					6224.35	11748	2649.65		1676.74
P30740	P30740	ILEU_HUMAN	SERPINB1	7618.81	5375.48	8589.91	5786.22	3958.2	5113.11	13477	9814.79	14916.5
P30837	P30837	AL1B1_HUMAN	ALDH1B1	7401.75	5489.12	7571.49	10856.6	12915.1	6594.3	11715.6	13792	12962
P30838	P30838	AL3A1_HUMAN	ALDH3A1		24450.2					1368.35		
P30876	P30876	RPB2_HUMAN	POLR2B	6301.61	6623.33	5913.48	8283.42	11298	11058.5	13628.7	10568.5	12878.8
P31025	P31025	LCN1_HUMAN	LCN1	33368.3		31319.1	25278.5		4546.81	6410.3	98664.8	5488.18
P31040	P31040	SDHA_HUMAN	SDHA			3727.95			1319.69	2810.34	2163.66	3503.31
P31150	P31150	GDIA_HUMAN	GDI1			1925.18	1686.51		795.825			561.3
P31151	P31151;Q86SG5	S10A7_HUMAN	S100A7	35884.2	31794.5	57931.3	34365.8	15572.2	20424.1	14924.7	46325.5	23886.1
P31153	P31153	METK2_HUMAN	MAT2A	9834.3	9644.68	8608.83	9347.17	6744.73	13370.3	28664.7	23303.3	29162.7
P31323	P13861;P31323	KAP3_HUMAN	PRKAR2B	7832.81	12944.7	5878.97	3143.2	2511.55	3068.13	2114	1067.8	1855.99
P31327	P31327	CPSM_HUMAN	CPS1									
P31350	P31350	RIR2_HUMAN	RRM2					1256.26	2119.96	1796.87	3777.93	2508.5
P31689	P31689	DNJA1_HUMAN	DNAJA1	1785.42	1105.93	6434.32	18204.2	17642.2	18045.4	27955.6	35422.3	26592.1
P31930	P31930	QCR1_HUMAN	UQCRC1	4696.55	4060.47	4332.45	2035.95	3180.09	3137.06	9383.1	5161.24	9350.27
P31937	P31937	3HIDH_HUMAN	HIBADH				1068.38	4124.73	9049.1	1180.32		1059.62
P31939	P31939	PUR9_HUMAN	ATIC	3376.11		5492.2	5918.77	7143.35	14019.8	5349.78	2357.3	5880.15
P31942	P31942	HNRH3_HUMAN	HNRNPH3					34513.7	7769.78	2606.15		6942.52
P31943	P31943	HNRH1_HUMAN	HNRNPH1	35072.8	38612	37958.4	32212.4	82975.7	34858.3	34170.3	23934.8	36214.1
P31944	P31944	CASPE_HUMAN	CASP14	18282.4	13766.4	22697.4	12292.7	12767.8	5891.15	6646.26	10946.6	9757.36
P31946	P31946	1433B_HUMAN	YWHAB			13198.7	5361.87	4978	11048.8	6559.11	3634.9	5933.92
P31947	P31947	1433S_HUMAN	SFN	6725.88	4383.89	5282.95	3538.05	2587.92	6620.09	4698.8	2492.72	5095.51
P31948	P31948	STIP1_HUMAN	STIP1	9053.8	3157.57	9153.4	17504.8	21073	39309.2	18992.2	7518.16	18299.4
P31949	P31949	S10AB_HUMAN	S100A11	7128.61			8456.78	5949.04	9587.55	6600.2	7485.55	4630.28
P32004	P32004	L1CAM_HUMAN	L1CAM					1894.28	1031.3	1773.93		1432.12
P32119	P32119	PRDX2_HUMAN	PRDX2	12738.4	9016.18	16750.2	7635.6	7089.77	8353.54	18992.7	8373.54	16555.2
P32189;Q 14409	Q14410;P32189;Q14409	GLPK3_HUMAN;GLPK_HUMAN	GK;GK3	14416.8	12519.3	7982.1						
P32320	P32320	CDD_HUMAN	CDA	4312.34	3102.43	5635.98	3229.08		2434.72	2305.23		2936.58
P32322	P32322	P5CR1_HUMAN	PYCR1	2855.94	4039.22	3931.39	3804.24	4556.4	5142.96	10620.8	12857.6	15637.2
P32455	P32455	GBP1_HUMAN	GBP1				1499.59	1822.33	1667.97	1533.8	1167.26	5693.56
P32929	P32929	CGL_HUMAN	CTH					1011.47	1463.91			879.618
P32969	P32969	RL9_HUMAN	RPL9P9	16331	12248.5	19346.4	15528.4	29555.2	19848.5	15296.7	16090.6	14971.1
P32970	P32970	CD70_HUMAN	CD70				3569.96	6635.94	6977.83			
P33121	P33121	ACSL1_HUMAN	ACSL1	3469.34	4082.41	5615.9				3937.76	3110.66	4058.07
P33176	P33176	KINH_HUMAN	KIF5B	10808.9	5524.94	9877.29	11052.3	9683.35	9016.5	7308.52	6685.45	6192.49
P33316	P33316	DUT_HUMAN	DUT			9159.15	6226.09	5008.37	10316.7	9032.09	6058.49	7560.99
P33527	P33527	MRP1_HUMAN	ABCC1		1859.57				614.019	1698.8	594.481	610.663
P33897	P33897	ABCD1_HUMAN	ABCD1							2633.98		1762.21
P33981	P33981	TTK_HUMAN	TTK				775.77	1384.19		949.697	1436.09	1081.76
P33991	P33991	MCM4_HUMAN	MCM4	7179.63	9351.54	10993.3	12268.4	17869.7	14083.3	20961.8	15358	18922.5
P33992	P33992	MCM5_HUMAN	MCM5	15379.8	15627.3	13321.2	6675.02	4692.32	7192.3	9765	6125.75	9247.51
P33993	P33993	MCM7_HUMAN	MCM7	18177.4	21006.8	14567.7	15426.4	23236.3	18115.3	22743.2	18761.7	24218.7
P34897	P34897	GLYM_HUMAN	SHMT2	9441.25	7434.3		13680.3	14004.1	16116.1	14033.3	13845.5	11120.5
P34932	P34932	HSP74_HUMAN	HSPA4	7709.31	5207.55	7555.27	9792.98	13054.3	16519.1	8736.13	4885.63	8771.32
P35030	P35030	TRY3_HUMAN	PRSS3	3679.55	860.946	2958.46	5174.83	1052.67	1115.03	2602.09	8053.92	1682.35
P35080	P35080	PROF2_HUMAN	PFN2	9666.93		6086.71	6306.48	1739.23	950.741	1713.42	5063.99	
P35221	P35221	CTNA1_HUMAN	CTNNA1		2546.49	3936.08	6182.5	11914.1	9494.13	14828.4	5590.91	12198.6
P35222	P35222	CTNB1_HUMAN	CTNNB1				3624.1	6143.65	3026.03	8319.53	4191.69	7696.97
P35232	P35232	PHB1_HUMAN	PHB1	5641.57		6968.86	5081.43	7968.21	4229.23	9191.85	7863.36	11070.9
P35237	P35237	SPB6_HUMAN	SERPINB6	9980.42	3853.53	12716	10911	6885.21	13898.5	15877.8	24783.7	19794
P35241	P35241	RADI_HUMAN	RDX	6846.81		4688.61	3839.72	6295.98	9331.15	3759.87		2771.77
P35244	P35244	RFA3_HUMAN	RPA3	85803.7	335084	5822.2	26005.1	214681	21010.4	14653.4	81210.3	2895.37
P35249	P35249	RFC4_HUMAN	RFC4	4329.49	5494.51	4097.48	7756.42	9087.54	6788.17	16372.8	13338.6	18088
P35250	P35250	RFC2_HUMAN	RFC2				4568.31	4955.75	2920.56	6868.75	7908.55	8295.77
P35251	P35251	RFC1_HUMAN	RFC1		1984.73			3197.05	1336.94	1485.31	1238.61	1816.41
P35268	P35268	RL22_HUMAN	RPL22	33297.3	22531.1	25015.3	27641.2	63677.5	29617.1	24597.7	42928.3	26298.8
P35269	P35269	T2FA_HUMAN	GTF2F1				1064.44	1192.17	704.227	486.976	1112.18	617.883
P35270	P35270	SPRE_HUMAN	SPR						2572.87	1139.59		1668.02
P35520	P35520	CBS_HUMAN	CBS				3081.09	5203.13	7241.47	1737.11		
P35527	P35527	K1C9_HUMAN	KRT9	#########	#########	2.19672E+06	1.35603E+06	377223	504654	488492	1.21211E+06	858432
P35568	P35568	IRS1_HUMAN	IRS1	1583.7				1851.61				
P35573	P35573	GDE_HUMAN	AGL				954.676	13032.2	7400.67	1199.3	1613.37	2848.89
P35579	P35579	MYH9_HUMAN	MYH9	117151	139364	80665.3	404898	733214	384457	222196	192565	157708
P35580	P35580;Q7Z406	MYH10_HUMAN	MYH10	28393.1	31956.6	20729.8	115533	207347	106216	141080	117382	104737
P35606	P35606	COPB2_HUMAN	COPB2	7087.58	6853.68	9283.15	12005.5	15051.4	14856.2	14073.4	12435	12158.9
P35609	P35609	ACTN2_HUMAN	ACTN2									
P35610	P35610	SOAT1_HUMAN	SOAT1		1395.47			2965.12	1895.99	773.26		989.47
P35611	P35611	ADDA_HUMAN	ADD1				2270.41	1910.51	1550.15	1397.53	1822.15	2071.15
P35613	P35613	BASI_HUMAN	BSG	16056.6	6224.84	12270.4	12595.2	7612.1	8523.6	60236.6	39587.8	35084.4
P35637	P35637	FUS_HUMAN	FUS	6857.2	7856.92	6834.84	5159.55	30497.1	14241	4956.26	6099.94	4889.2
P35658	P35658	NU214_HUMAN	NUP214				2209.6	1690.87	1608.76	1408.77	1955.08	1866.44
P35659	P35659	DEK_HUMAN	DEK	9092.58	9697.5	8510.08	6258.61	15371	5002.85	4565.42	6925.34	6117.32
P35754	P35754	GLRX1_HUMAN	GLRX	9736.48	4827.97	9157.14	6119.33	7824.03	8843.41	6563.17	4216.83	7729.5
P35869	P35869	AHR_HUMAN	AHR									
P35900	P35900	K1C20_HUMAN	KRT20				1486.57			772.959		523.862
P35908	P35908	K22E_HUMAN	KRT2	727684	667375	1.36308E+06	717011	226320	402435	559016	656337	803706
P35998	P35998	PRS7_HUMAN	PSMC2	19721.4	17559	20633.7	21607.1	20580	19815	16832	13275.2	15165.2
P36404	P36404	ARL2_HUMAN	ARL 2:00	3506.07			5217.15		2885.68			1467.68
P36507	Q02750;P36507	MP2K2_HUMAN	MAP2K2			9835.17	8790.07	5635.07	7752.43	9663.38	8709.84	9077.19
P36542	P36542	ATPG_HUMAN	ATP5F1C	20628.9	20653.8	24344.6	6873.44	9040.04	9567.48	24364	11189.4	21378.5
P36551	P36551	HEM6_HUMAN	CPOX	3723.63		3590.89	3735.29	4741.5	9514.74	2504.07		3347.64
P36578	P36578	RL4_HUMAN	RPL4	88321.4	88190.2	71958.3	137222	145789	73153.8	96435.9	91184.5	91898.9
P36639	P36639	8ODP_HUMAN	NUDT1				631.412	393.689	1163.79	532.066		402.019
P36776	P36776	LONM_HUMAN	LONP1	3672.13		4996.97	4247.93	4844.39	5957.98	4181.29	2434.26	5213.41
P36871	P36871	PGM1_HUMAN	PGM1			1373.45			2305.69	794.259		
P36915	P36915	GNL1_HUMAN	GNL1				1701.43	1756.94	3642.33	1082.54		
P36957	P36957	ODO2_HUMAN	DLST	14751.4	11290.6	22608.3	31162.2	4182.07	16156	13851.2	13774.6	18841.9
P37059	P37059	DHB2_HUMAN	HSD17B2	6281.56	6225.69	6184.9						
P37108	P37108	SRP14_HUMAN	SRP14	4573.91		7788.18	6110.83	10870.2	11390.1	8658.55	7408.19	5694.37
P37235	P37235	HPCL1_HUMAN	HPCAL1				1709.9	3978.02	5763.15	11119.7	2617.69	14783.1
P37268	P37268	FDFT_HUMAN	FDFT1				8894.13	3735.73	3626.37	2675.42	8455.93	3298.38
P37802	P37802	TAGL2_HUMAN	TAGLN2	20770.1	18426.4	19151.8	19394.9	7390.73	25416.8	8853.47	11685.6	9895.28
P37837	P37837	TALDO_HUMAN	TALDO1	14353.2		16370.1	5127.23	3692.23	8755.69	7753.51	3401.14	7327.11
P38117	P38117	ETFB_HUMAN	ETFB	3611.35	1644.1	3388.77	2229.63	3001.97	5187.56	5548.65	2491.74	4487.96
P38159	P38159	RBMX_HUMAN	RBMX	9310.26	4091.04	11545.2	7137.74	23698.5	10580	10031.2	7358.76	10808.3
P38606	P38606	VATA_HUMAN	ATP6V1A	6035.21	4467.5	5174.22	6585.79	5827.44	6156.6	3935.61	6119.07	5061.21
P38646	P38646	GRP75_HUMAN	HSPA9	110938	96596.4	83370.3	75698.9	100449	116234	74380.9	48289.6	73377.1
P38919	P38919	IF4A3_HUMAN	EIF4A3			13109.1	5838.06	6004.93	6039.25	7405.9	4063.26	7159.11
P39019	P39019	RS19_HUMAN	RPS19	118884	154498	114919	73561.5	155808	95765.1	104240	86957.8	85739.4
P39023	P39023	RL3_HUMAN	RPL3	62606.9	78027.4	48269.6	97938.5	152433	90895.1	83380.3	80437.9	71895.6
P39060	P39060	COIA1_HUMAN	COL18A1	14074.8	9689.03	12607.1			1888.33	6221.54	3558.6	7197.42
P39656	P39656	OST48_HUMAN	DDOST	22444.1	22011.7	20153.8	15750.3	16435	12420.2	36179.7	26309.5	23738.9
P39687	P39687	AN32A_HUMAN	ANP32A	5268.62		5629.41	4090.08	4208.48	10222	5282.3	5391.88	5048.8
P39748	P39748	FEN1_HUMAN	FEN1	5977.06			5880.18	9339.04	17093.9	9444.62	4442.98	6987.22
P39880	P39880	CUX1_HUMAN	CUX1	1999.45	8522.5			4135.48	2381.99	6634.8	4615.72	
P40121	P40121	CAPG_HUMAN	CAPG			2191.96	5345.35	2168.12	6527.99	1874.34		2830.9
P40123	P40123	CAP2_HUMAN	CAP2	5945.25	5632.61		12237.1	3214.66	8160.59	3241.38	15809.4	3771.87
P40199	P40199	CEAM6_HUMAN	CEACAM6							5462.87	2119.72	4730.4
P40227	P40227	TCPZ_HUMAN	CCT6A	15282.1	8485.02	17395.4	17728.1	19654.5	27061.7	36646.3	16226.5	37782.9
P40425	P40425	PBX2_HUMAN	PBX2					6114.94	3305.89	1426.39		1014.78
P40429	Q6NVV1;P40429	RL13A_HUMAN	RPL13A	11452.7	13724.3	12495	13091.9	17016.5	16857.4	13233.6	12227.4	13627.6
P40616	P40616	ARL1_HUMAN	ARL 1:00	3744.89		9722.17	10316.5		1748.85	3253.23	8794	5773.55
P40692	P40692	MLH1_HUMAN	MLH1				900.866	845.026	919.552	942.142	1344.36	910.099
P40763	P40763	STAT3_HUMAN	STAT3	3268.61	3689.96	3871.87	5095.83	4484.85	4478.52	2620.6	1181.35	2379.3
P40818	P40818	UBP8_HUMAN	USP8	1593.21	2265.46	1420.81	4053.24			1179.32		1251.54
P40925	P40925	MDHC_HUMAN	MDH1	10925.8	14261.8	13492.6	12872.4	4180.96	19046.4	9371.37	6767.76	9888.93
P40926	P40926	MDHM_HUMAN	MDH2	16938.5	7228.08	16000.5	12855.4	8245.42	17088.8	12487	14828.2	11476.9
P40937	P40937	RFC5_HUMAN	RFC5	3762.03	3897.59	4366.18	8662.88	5328.67	4852.07	15217.5	18337.6	18812.7
P40938	P40938	RFC3_HUMAN	RFC3	3422.79	4955.41	2276.5	5515.95	5554.44	4961.69	9026.47	7370.24	9575.21
P40939	P40939	ECHA_HUMAN	HADHA	7790.18	5800.15	6636.04	6666.98	5736.06	5729.83	8061.77	7215.35	8148.15
P41091	Q2VIR3;P41091	IF2G_HUMAN	EIF2S3	4292.77	3764.89	3778.77	9799.13	13327.8	15820.6	10926	9268.58	10281.6
P41162	P41162	ETV3_HUMAN	ETV3				3711.65	1917.1	2401.8	5153.34	4426.54	3570.94
P41208	Q12798;P41208	CETN2_HUMAN	CETN2		12311			5114.81				
P41227	P41227	NAA10_HUMAN	NAA10					9394.65	6259.33	9609.18	5218.05	8271.92
P41240	P41240	CSK_HUMAN	CSK		1743.6	1458.46		10510.6	11421.9	8279.99	2166.53	9538.99
P41247	P41247	PLPL4_HUMAN	PNPLA4				201.472	999.353	478.625	924.27		1470.09
P41250	P41250	GARS_HUMAN	GARS1	7411.66	4208.72	6577.57	8830.38	8988.24	14414.1	5321.5	3058.45	5699.76
P41252	P41252	SYIC_HUMAN	IARS1	15380.4	17826.2	19030.9	34562.2	21220.2	22999.5	17607.9	22091	19004.9
P41743	P41743	KPCI_HUMAN	PRKCI				2629.8	5441.87	6611.77	3180.03	2951.6	2942.48
P42025	P42025	ACTY_HUMAN	ACTR1B			1737.91		1361.12	2187.7	632.544		
P42126	P42126	ECI1_HUMAN	ECI1	3349.17		4451.75	2870.74		5621.18	4649.76	3908.97	5221.34
P42166	P42166	LAP2A_HUMAN	TMPO	19183.8	16022.3	15666.9	31020.4	32632.4	20780.5	10629.9	18870.4	13666.1
P42224	P42224	STAT1_HUMAN	STAT1	32279.6	37419.1	38598.1	13290.3	9674.36	12641.9	9560.4	9535.98	9858.8
P42226	P42226	STAT6_HUMAN	STAT6	2958.04	3756.29	3222.91	4526.39	2944.86	4265.14	2034.92	1847.57	2655.15
P42229	P42229	STA5A_HUMAN	STAT5A									
P42285	P42285	MTREX_HUMAN	MTREX				1132.4	6769.6	3834.74	3036.87	2624.22	3317.38
P42330	P42330	AK1C3_HUMAN	AKR1C3	3914.77		3216.82				2024.97		1733.7
P42345	P42345	MTOR_HUMAN	MTOR	3545.54	2757.66	3274.3	4291.07	4295.57	4393.66	3265.35	7458.98	4744.38
P42356	P42356	PI4KA_HUMAN	PI4KA			1644.54	2923.72	5967.17	5627.39	5645.12	2273.27	5164.35
P42357	P42357	HUTH_HUMAN	HAL			5394.42	5620.35		2865.89	1455.5	5683.49	2347.3
P42566	P42566	EPS15_HUMAN	EPS15		457.092	1697.82	2279.44	4999.29	4672.45	2187.62		1532.6
P42677	P42677;Q71UM5	RS27_HUMAN	RPS27	25966.1	31585.1	27126.6	35805.4	23965.4	25637.5	39703.7	45205.2	37467.5
P42694	P42694	HELZ_HUMAN	HELZ		770.123			5367.51	2223.53	1359.5	495.453	
P42695	P42695	CNDD3_HUMAN	NCAPD3				587.989			866.704	1733.32	1442.75
P42704	P42704	LPPRC_HUMAN	LRPPRC	8098.65	11136.4	10497.1	13331.3	10477.1	12016.3	18205.1	16127.2	18540
P42765	P42765	THIM_HUMAN	ACAA2	4489.08		4617.37	1885.57	1779.74	4067.89	2645.03	2172.63	3046.51
P42766	P42766	RL35_HUMAN	RPL35	19687.5	19301.4	11735.3	20816.2	49401.3	18690.6	16749.8	19430.5	13819
P42771	P42771	CDN2A_HUMAN	CDKN2A	4337.01	2353.39	4372.15			802.588			
P42785	P42785	PCP_HUMAN	PRCP				720.911	3237.65	5037.24	4402.4	1053.89	2810.72
P42892	P42892	ECE1_HUMAN	ECE1	3304.54		1779.66	1843.71	1781.12	2069.57	895.903	5208.48	1746.39
P43034	P43034	LIS1_HUMAN	PAFAH1B1	2478.87		2705.33	1788.3	2363.19	4400.69	3932.04	2585.95	3481.97
P43121	P43121	MUC18_HUMAN	MCAM				15989.5	34543	33033.9	6327.06	4393.95	7609.73
P43243	P43243	MATR3_HUMAN	MATR3	6534.96	8717.22	6949.09	5247.78	6544.73	2706.11	4523.93	5347.27	4778.3
P43246	P43246	MSH2_HUMAN	MSH2	19622.5	28279.7	6623.34	18774.1	16536.5	8445.61	9695.02	18775.5	10591.6
P43250	P43250	GRK6_HUMAN	GRK6							2226.36		2962.48
P43304	P43304	GPDM_HUMAN	GPD2	5708.45	3491.55	5856.71	3007.11	3015.03	2351.02	2944.62	2481.87	2997.44
P43307	P43307	SSRA_HUMAN	SSR1					2267.68	2448.37	1158.61	1313.47	3656.75
P43362	P43362	MAGA9_HUMAN	MAGEA9B				2600.15	2011.33	1952.08			
P43405	P43405	KSYK_HUMAN	SYK							8449.8	6961.28	8427.59
P43487	P43487	RANG_HUMAN	RANBP1				3157.41	1864.3	3855.86	2446.24		2682.78
P43490	P43490	NAMPT_HUMAN	NAMPT	18941.1	11983	22578.2	9858.76	11552.6	31120.6	7146.17	4297.72	8495.22
P43686	P43686	PRS6B_HUMAN	PSMC4	19698.4	16756.1	20811.1	22065.1	22723	24508.6	22714	20074.4	22519
P43699	P52952;Q8H2Z4;P43699	NKX21_HUMAN	NKX2-1			4345.05			1498.58			
P45880	P45880	VDAC2_HUMAN	VDAC2	21174.1	16794	19616.9	3988.55	4260.47	4397.77	8101.52	7861.11	7993.29
P45954	P45954	ACDSB_HUMAN	ACADSB	2968.95		3328.8	1294.8	1839.4	1525.55	3150.45	5118.95	3776.71
P45974	P45974	UBP5_HUMAN	USP5	3048.02		7112.44	6524.11	5658.73	10835.9	6851.53	3295.27	6566.75
P46013	P46013	KI67_HUMAN	MKI67							2497.11		2369.84
P46020	P46020	KPB1_HUMAN	PHKA1	15786.6				5132.84	3773.41	3525.21		3690.11
P46060	P46060	RAGP1_HUMAN	RANGAP1	4201.47		5393.01	6733.39	4163.57	5026.23	3970.93	4683.54	3801.37
P46063	P46063	RECQ1_HUMAN	RECQL	15419.8	24096.4	3289.58	12227.8	13461.6	7620.52	3971.3	18113.8	4779.09
P46087	P46087	NOP2_HUMAN	NOP2	5631.51	4146.86	4349.22	1556.82		1652.9	3801.12	5324	5049.59
P46109	P46109	CRKL_HUMAN	CRKL					2078.55	3343			
P46379	P46379	BAG6_HUMAN	BAG6	6784.69	6805.28	8419.38	6808.06	8052.55	10912.9	7264.55	4614.76	8242.7
P46439	P46439	GSTM5_HUMAN	GSTM5				3677.81					
P46459	P46459	NSF_HUMAN	NSF	18635.9	16041.2	17606.5	22003.8	12094	12203.4	12795.7	17316.9	13928.8
P46734	P46734	MP2K3_HUMAN	MAP2K3			1989.8	2785.35	3162.9	2084.45	1042.9	2726.84	2051.33
P46776	P46776	RL27A_HUMAN	RPL27A	58575.1	53500.8	58557	83437.6	80556.5	45411.8	44165.1	112346	54588.5
P46777	P46777	RL5_HUMAN	RPL5	22809.3	19501	29459.6	41072.1	66531.6	92338.8	52335.2	30346.9	45398.4
P46778	P46778	RL21_HUMAN	RPL21	9488.2	10048.6	11088.3	13832.4	18371.2	17921.2	15120.6	11146.7	17739
P46779	P46779	RL28_HUMAN	RPL28	25279	18689.6	24821.8	37564.9	22867.1	20085.8	21135	36442.9	25218
P46781	P46781	RS9_HUMAN	RPS9	73581.6	78801.9	69792.1	84839.4	127985	95691.1	87952.9	64496.4	97564.5
P46782	P46782	RS5_HUMAN	RPS5	23852.8	35580.1	30451.5	26079.1	48807.6	40690.4	34552.9	28171.4	34188.8
P46783	Q9NQ39;P46783	RS10_HUMAN	RPS10	36276.9	38677.6	36766.8	42789.3	122975	67059.6	70127	40853.6	65507.9
P46821	P46821	MAP1B_HUMAN	MAP1B		13735.2	3725.65	11180.5	10766.8	12863.9			
P46926	P46926;Q8TDQ7	GNPI1_HUMAN	GNPDA1			3025.59			3216.57	1296.57		
P46934	P46934	NEDD4_HUMAN	NEDD4	945.653	895.375	1363.97	1222.29	1471.31	2572.38	807.122		665.208
P46937	P46937	YAP1_HUMAN	YAP1					240.877	621.761			
P46940	P46940	IQGA1_HUMAN	IQGAP1	13975.2	12708.7	12001.6	19673	13910.9	13681.1	8699.54	7685.92	9025.78
P46977	P46977	STT3A_HUMAN	STT3A	6834.48	9717.11	7820.21	6362.09	6345.86	6431.72	6030.06	5077.03	6535.46
P47755	P47755	CAZA2_HUMAN	CAPZA2	4668.27	4651.98	3274.52	9396.3	26417.4	20367.9	8054.71	2167.84	9937.57
P47756	P47756	CAPZB_HUMAN	CAPZB	17186.5	13701.9	23323	27265.1	57405.7	47097.9	39251	17817	31416
P47813	P47813	IF1AX_HUMAN	EIF1AX				1559.84	1560.56	2658.79	1519.55	2378.89	1407.23
P47895	P47895	AL1A3_HUMAN	ALDH1A3		5774.78	5006.27		4529.44		6337.48	7156.33	5301.64
P47897	P47897	SYQ_HUMAN	QARS1	7536.04	6294.77	7838.28	12342.7	7323.16	8958.08	4615.61	5170.85	6029.04
P47914	P47914	RL29_HUMAN	RPL29	10213.6	6674.52	5785.59	13059.1	8070.98	4149.56	5077.44	14949.8	4377.49
P47929	P47929	LEG7_HUMAN	LGALS7B	3268.19		5014.86	2944.94	3499.2	3682.72	2736.26	3858.97	2564.67
P47974	P47974	TISD_HUMAN	ZFP36L2	7246.77	7999.42	6735.61	3415.63	4995.46	3211.27	8322.51	5536.24	10882.6
P48047	P48047	ATPO_HUMAN	ATP5PO	9097.92	6873.38	7306.15	3711.25		2671.81	5080.47	4168.67	3609.47
P48147	P48147	PPCE_HUMAN	PREP				2409.24	4520.1	6244.35	2181.99		3165.55
P48163	P48163	MAOX_HUMAN	ME1									
P48444	P48444	COPD_HUMAN	ARCN1	10054.4	8345.38	9565.03	17239.5	20621.2	23190.8	15563.3	14372.5	12867.6
P48506	P48506	GSH1_HUMAN	GCLC						3315.91	2184.99		1438.36
P48507	P48507	GSH0_HUMAN	GCLM		1152.27	2080.39		836.267	2659.83	833.914		835.483
P48556	P48556	PSMD8_HUMAN	PSMD8	7506.55	5640.27	6730.47	8501.11	4206.26	6537.12	5734.36	7392.45	8328.61
P48594	P48594	SPB4_HUMAN	SERPINB4	3425.97		5932.74	2970.31		2526.9	1734.34	5193.51	4754.94
P48634	P48634	PRC2A_HUMAN	PRRC2A				1841.72	2684.48	1687.38	886.101		688.301
P48637	P48637	GSHB_HUMAN	GSS			1647.58	1484.14	1638.43	2843.23	1790.1	4758.13	2593.19
P48643	P48643	TCPE_HUMAN	CCT5	10232.7	9280.62	11184.5	14517.8	21140.6	29672.6	25121.6	10771.4	28695.8
P48651	P48651	PTSS1_HUMAN	PTDSS1							1494.6	2470.93	2747.53
P48668	P48668	K2C6C_HUMAN	KRT6C			8744.69					8904.51	5378.1
P48681	P48681	NEST_HUMAN	NES	26967.1	31086.8	23850.8	5603.3	11883.1	9773.42	84814	11780.1	83471.9
P48723	P48723	HSP13_HUMAN	HSPA13			574.142	1249.05	1168.38	2213.44	656.331		
P48729	P48729;Q8N752	KC1A_HUMAN	CSNK1A1	3739.95	5976.04	7092.3	3858.69	18608.1	14916.8	25058.9	3431.75	27282.8
P48735	P48735	IDHP_HUMAN	IDH2	6658.18	8677.16	9391.19	5589.35	4894.99	6221.03	13749.3	16352.7	14938.3
P48739	P48739	PIPNB_HUMAN	PITPNB			3428.37	2363.33	2086.49	6761.55	2602.26	1241.02	2394.78
P48960	P48960	AGRE5_HUMAN	ADGRE5		7650.94	4942.2	7237.11	4885.89	5771.75	7083.94	6403.23	5460.05
P49005	P49005	DPOD2_HUMAN	POLD2		1878.1		1377.92	4558.99	3435.74	4498.47	2535.92	4695.84
P49006	P49006	MRP_HUMAN	MARCKSL1				12818.9	6784.4	10776.6	137871	19393.9	108248
P49023	P49023	PAXI_HUMAN	PXN	4017.06		2864.46	2915.28		1306.76	1694.2	2199.13	1760.07
P49184	P49184	DNSL1_HUMAN	DNASE1L1				1571.18			5301.56	1185.97	4881.14
P49189	P49189	AL9A1_HUMAN	ALDH9A1	10946.6		11491.7	6866.49	5580.42	8869.21	6121.96	11366.9	6072.49
P49207	P49207	RL34_HUMAN	RPL34	11023.4	12093.9	6873.13	7688.84	18488	11096	9592.79	8944.39	9862.55
P49247	P49247	RPIA_HUMAN	RPIA				628.02	967.94	1598.1	1024.66	446.025	806.049
P49257	P49257	LMAN1_HUMAN	LMAN1	4988.46	4341.89	3005.56	2683.77	6168.79	6053.49	6076.11	1397.52	4922.11
P49321	P49321	NASP_HUMAN	NASP	3512.97	1478.86	3205.98	4031.35	3986.36	11370.8	4841.65	1508.45	4637.32
P49327	P49327	FAS_HUMAN	FASN	9469.87	6958.51	9367.78	14488.9	10832	11368.9	18760.6	14696.6	16982.6
P49368	P49368	TCPG_HUMAN	CCT3	14677.5	9561.03	14615.3	30150.7	23534.6	38540	39347	30895.3	39817.7
P49406	P49406	RM19_HUMAN	MRPL19			2636.33				2664.58		3486.33
P49411	P49411	EFTU_HUMAN	TUFM	40508.8	35469.6	40663	68844.1	54706.6	54357	49704.8	59768.1	50716.4
P49419	P49419	AL7A1_HUMAN	ALDH7A1	11008.8	6561.83	9401.49	5636.33	8016.43	8340.88	6872.37	5470.6	6838.7
P49458	P49458	SRP09_HUMAN	SRP9			3130.76	3636.5	2627.36	2072.41	2860.87	3490.6	2228.56
P49585	P49585;Q9Y5K3	PCY1A_HUMAN	PCYT1A					2443.95		1566.85		
P49588	P49588	SYAC_HUMAN	AARS1	6945.04	1785.89	5155.2	5315.15	5812.6	6872.52	6564.3	4644.23	5429.7
P49589	P49589	SYCC_HUMAN	CARS1	14511.6	12886.2	14725.8	12659.1	15271.7	12258.1	6783.13	7428.45	6655.9
P49591	P49591	SYSC_HUMAN	SARS1	5223.21		5640.79	6768.27	11065.5	14554.8	6881.01	2683.49	4804.64
P49593	P49593	PPM1F_HUMAN	PPM1F	1672.29		2043.32	1018.22		1545.31	1036.53		1021.46
P49642	P49642	PRI1_HUMAN	PRIM1				2358.22	2546.48	2641.86	4146.99	3674.79	3076.58
P49643	P49643	PRI2_HUMAN	PRIM2					1898.07	1163.24	2258.86		5020.74
P49720	P49720	PSB3_HUMAN	PSMB3	2718.12	1246.32	3981.41	1013.18	1816.46	4478.99	2763.89	1206.84	4285.06
P49721	P49721	PSB2_HUMAN	PSMB2			6609.93	3881.32	12614	15703.4	7955.84	4619.76	10775.8
P49736	P49736	MCM2_HUMAN	MCM2			8546.74	1977.97	3698.16	5193.33	6724.28	3423.05	7047.68
P49748	P49748	ACADV_HUMAN	ACADVL	10188		8972.8	5925.43	6040.22	7814.69	8428.45	11963.9	9428.57
P49750	P49750	YLPM1_HUMAN	YLPM1					2250.08				
P49755	P49755	TMEDA_HUMAN	TMED10		11501.8	14125.1	4182.03	6707.15	8047.74	6384.81	5580.09	8195.52
P49756	P49756	RBM25_HUMAN	RBM25	3352.92	3828.27	3261.44	3413	3128.54	2261.53	3845.4	4595.74	3063.38
P49770	P49770	EI2BB_HUMAN	EIF2B2		991.493		3162.02	2163.86	2227.68	2152.59	3212.94	2321.48
P49773	P49773	HINT1_HUMAN	HINT1						1206.89		857.479	
P49790	P49790	NU153_HUMAN	NUP153				1435.81	2803.04	2012.39	6081.15		
P49792	A6NKT7;O14715;Q7Z3J3;P0DJD1;P49792;P0DJD0;O	RBP2_HUMAN	RANBP2	2756.47		2394.43	4897.08	3339.69	4082.25	5197.24	6389.7	5927.68
P49795	P49795	RGS19_HUMAN	RGS19					6935.11	5071.97	9000.49		8074.57
P49815	P49815	TSC2_HUMAN	TSC2	2703.44	2050.71	2831.61	2899.15	2315.91	1892.24	1569.82	1011.6	1531.73
P49821	P49821	NDUV1_HUMAN	NDUFV1					5488.1		2520.22	4264.7	1548.31
P49841	P49841	GSK3B_HUMAN	GSK3B				898.404				1680.61	381.015
P49902	P49902	5NTC_HUMAN	NT5C2	4257.33		6969.67	4542.15	2633.92	4536.18	2091.67	1944.45	3185.37
P49903	P49903	SPS1_HUMAN	SEPHS1				1084.6		2993.14	1167.72	848.567	439.526
P49915	P49915	GUAA_HUMAN	GMPS	5443.73	3917.15	6106.5	9281.97	13680.7	14254.1	7701.08	6832.65	7845.88
P49916	P49916	DNLI3_HUMAN	LIG3	35037.4	69113.6	3795.21	58587.5	98827.7	9051.92	6742.27	7510.46	5328.41
P49917	P49917	DNLI4_HUMAN	LIG4				1429.35		1332.88	1384.32		
P49959	P49959	MRE11_HUMAN	MRE11			3520.9	4572.75	6075.49	1813.99	3475.41	4585.66	3144.56
P50148	P50148	GNAQ_HUMAN	GNAQ				9352.4	4606.32	5742.77	14093.4	7620.1	12154.7
P50150	P50150	GBG4_HUMAN	GNG4						2684.15	34601.8	5676.58	29676.8
P50151	P50151	GBG10_HUMAN	GNG10							2791.41		5352.28
P50213	P50213	IDH3A_HUMAN	IDH3A	4155.31	3515.7	4604.1	12295.3	5037.27	8010.38	14298.4	13379.6	15369.9
P50395	P50395	GDIB_HUMAN	GDI2	11125.4	5202.03	13396.1	9556.05	7617.58	13744	10252.1	7634.77	8219.54
P50402	P50402	EMD_HUMAN	EMD	13735.4	15894.7	24611.5	6159.72	9882.86	9266.76	13164.8	12533.5	13761.5
P50416	P50416	CPT1A_HUMAN	CPT1A	4083.25	5446.63	3750.79				9200.77	7841.21	10895.5
P50452	P50452	SPB8_HUMAN	SERPINB8	4796.9		6927.47	3938.2				4451.63	5663.08
P50454	P50454	SERPH_HUMAN	SERPINH1	17169.6	8414.22	17162.7	12474.2	34748	56732.2	8265.07	4538.98	9109.89
P50502	Q8IZP2;P50502	F10A1_HUMAN	ST13			4967.03	8099.42	3629.91	6483.65	6820.18	5884.79	5655.5
P50552	P50552	VASP_HUMAN	VASP			1861.45	2360.5	1556.1	3726.28	1678.3		1990.67
P50570	Q9UQ16;P50570	DYN2_HUMAN	DNM2	7838.43	6466.5	8265.39	10782.3	8978.44	5662.29	13357.6	10918.2	11404.3
P50579	P50579	MAP2_HUMAN	METAP2			8006.6	3932.77	7295.63	7084.07	3088.38	3758.09	3626.04
P50747	P50747	BPL1_HUMAN	HLCS	7992.13	8745.34	11215.9	8238.11	6478.45	7096.69	10475.6	14293.1	10597
P50748	P50748	KNTC1_HUMAN	KNTC1				3455.84	5086.54	4330.69	4455.44	4261.34	5381.91
P50750	P50750	CDK9_HUMAN	CDK9				2628.91	1816.93	2877	4706.63	3454.51	3964.51
P50851	P50851	LRBA_HUMAN	LRBA				2847.75			5466.55	5490.29	4869.68
P50895	P50895	BCAM_HUMAN	BCAM				687.975	1947.64	1690.84	4353.53		3830.58
P50897	P50897	PPT1_HUMAN	PPT1			7697.46	14985.2	6108.1	6885.59	4821.16	14875.7	5159.28
P50914	P50914	RL14_HUMAN	RPL14	23218.9	21387.5	20717.7	24769.6	26987.1	26458.4	33843.5	18959.7	28676.6
P50990	P50990	TCPQ_HUMAN	CCT8	15911.3	5214.84	13938.5	31446.7	17868.5	20179.5	26507.6	21477.4	22810
P50991	P50991	TCPD_HUMAN	CCT4	8708.76	7511.88	10850.5	14246.2	12759.9	18206.9	16662.9	9568.32	15975.8
P50995	P50995	ANX11_HUMAN	ANXA11	9886.77	8659.22	9164.64	2559.27	7175.28	10275.7	8034.19	3293.83	7240.69
P51003	P51003	PAPOA_HUMAN	PAPOLA			1341.69		1742.1	2592.92	1344.24		528.571
P51114	P51114	FXR1_HUMAN	FXR1	9695.55	11639.6	6858.23	8373.24	19961.2	10425.4	4872.67	4149.01	5461.06
P51116	P51116	FXR2_HUMAN	FXR2	4672.71	7561.17	3335.72	2498.02	10715.1	3656.51	4255.69	2715.09	4407.98
P51148	P20339;P61020;P51148	RAB5C_HUMAN	RAB5C	19558.4	8370.93	9971.63	9592.94	19092.1	5378.62	12341.6	10062.6	8397.9
P51149	P51149	RAB7A_HUMAN	RAB7A	14978.7	12286.9	19035.5	12877.7	18904.1	22512.9	20797.9	13404.1	19382.2
P51151	P51151	RAB9A_HUMAN	RAB9A			4663.03	1598.99	2839.58	3558.55	4302.87	1497.57	3858.08
P51153	P51153	RAB13_HUMAN	RAB13				4474.63	3568.45	4749.77	6854.3	4516.82	5855
P51178	P51178	PLCD1_HUMAN	PLCD1				2893.06			2445.84		2101.87
P51398	P51398	RT29_HUMAN	DAP3				1609.95	2717.71	1244.48	2659.45	436.745	2721.05
P51532	Q6ZRS2;P51531;P51532	SMCA4_HUMAN	SMARCA4	4974.04	5403.61	3729.46	4106.53	7147.47	3850.2	4317.75	4083.39	4284.15
P51553	P51553	IDH3G_HUMAN	IDH3G	7542.18	8310.22	10927.5	4080.93	6152.04	6255.85	14443.4	9192.4	15524.3
P51570	P51570	GALK1_HUMAN	GALK1	12066.2	13238.5	8657.05	25039.4	5474.66	5808.72	6850.47	11143.5	6906.06
P51571	P51571	SSRD_HUMAN	SSR4	9837.7	9424.44	12542.3	7387.41	7038.28	4648.41	11968.2	9782.77	14662.9
P51572	P51572	BAP31_HUMAN	BCAP31	1764.06		2774.17	2184.84	4711.62	4130.34	5191.2	2486.19	5983.4
P51587	P51587	BRCA2_HUMAN	BRCA2			3169.21	2719.06	1072.55	1647.75	3250.99	2619.94	3200.03
P51608	P51608	MECP2_HUMAN	MECP2	4554.82		276.222	624.423					
P51610	P51610	HCFC1_HUMAN	HCFC1	2855.8	2173.45	3596.78	2383.82	2541.68	3057.2	4435.89	757.697	3455.86
P51617	P51617	IRAK1_HUMAN	IRAK1			1713.84	1705.05	2999.92	3124.11	3898.92	3096.95	4739.16
P51648	P51648	AL3A2_HUMAN	ALDH3A2			9678.65	2713.25	7696.64	3024.9	6083.75	6149.87	7363.89
P51649	P51649	SSDH_HUMAN	ALDH5A1							1529.82	2849.31	1426.63
P51659	P51659	DHB4_HUMAN	HSD17B4	13896	9542.23	10387.6	6806.34	17088.4	20585.9	10596.7	3405.38	9277.8
P51665	P51665	PSMD7_HUMAN	PSMD7	3076.06	4008.74	4533.57	9281.23	5184.03	8161.91	6527.21	4169.92	5733.66
P51688	P51688	SPHM_HUMAN	SGSH	9626.28		8863.21	4532.52			1831.54		1857.69
P51812	P51812	KS6A3_HUMAN	RPS6KA3	21865.7		10375.5	16524.4	25236.8	9766.58	15626.2	12459.4	11198.2
P51843	P51843	NR0B1_HUMAN	NR0B1									
P51858	P51858	HDGF_HUMAN	HDGF	5295.1		6668.24	5568.98	3011.69	8272.93	4641.85	3983.89	5327.01
P51970	P51970	NDUA8_HUMAN	NDUFA8				775.152	5575.34	3238.32	3823.15	3046.46	1585.14
P51991	P51991	ROA3_HUMAN	HNRNPA3	3185.02	1666.32		2783.24	10250.1	3594.74	1852.89	986.642	1872.88
P52209	P52209	6PGD_HUMAN	PGD	12415.4	5337.2	18936.9	10608.6	6104.33	12606.7	12127.2	10063.5	11606.6
P52272	P52272	HNRPM_HUMAN	HNRNPM	54788.1	42919	33227.2	43398.3	57788.4	23341.2	17936.8	18785	18984
P52292	P52292	IMA1_HUMAN	KPNA2	6481.66	5520.67	5969.11	28310.5	16269.1	19134.2	26226.2	24558.9	20858.9
P52294	P52294	IMA5_HUMAN	KPNA1	1868.95		1331.4	3493.62	4431.78	2880.18	3091.35	2453.02	2806.17
P52306	P52306	GDS1_HUMAN	RAP1GDS1			1933.6						
P52429	P52429	DGKE_HUMAN	DGKE							2104.4	1657.5	2149.79
P52434	P52434	RPAB3_HUMAN	POLR2H				2166.18		2450.44	2600.85		3805.21
P52564	P52564	MP2K6_HUMAN	MAP2K6	7333.59				3137.22	4054.19			
P52565	P52565	GDIR1_HUMAN	ARHGDIA	9670.17		16600.8	2993.26	11190.4	16145.5	7729.08	1927.35	10312.6
P52594	P52594	AGFG1_HUMAN	AGFG1	6352.88	7477.28	5501.85	3147.65		2170.24	2112.24	1131.59	2045.78
P52597	P52597	HNRPF_HUMAN	HNRNPF	1932.08	3877.95	4872.73	7410.48	16485.4	11141.3	9149.13	9859.14	6200.55
P52630	P52630	STAT2_HUMAN	STAT2		4583.16	2536.47	1373.3	8227.51	6946.23	3601.44		2179.74
P52701	P52701	MSH6_HUMAN	MSH6	4842.75	9362.61	6689.23	11693.1	13153.4	9256.27	12038.1	17987.8	11151.9
P52732	P52732	KIF11_HUMAN	KIF11				2222.14		1437.2	2990.9	2123.87	3345.84
P52789	P52789	HXK2_HUMAN	HK2	8075.21	7464.9	10966.6	13099.6	8275.8	10263.7	9946.03	9294.85	9011.25
P52815	P52815	RM12_HUMAN	MRPL12			3851.55	4353.17	4023.57	5380.01	7071.06	3644.03	6711.2
P52888	P52888	THOP1_HUMAN	THOP1						3355.53	2710.79		1982.16
P52895	Q04828;P52895	AK1C2_HUMAN	AKR1C2			6572.85						
P52907	P52907;P47755	CAZA1_HUMAN	CAPZA1	30024.6	29204.7	32372.5	67651.1	112426	99852.7	95711.8	35368	71646
P52926	P52926	HMGA2_HUMAN	HMGA2	22010.2		3379.84	82522	50111.6	43629.8	1665.56	16809.4	1712.13
P53004	P53004	BIEA_HUMAN	BLVRA				1748.62		6371.78			2106.42
P53041	P53041	PPP5_HUMAN	PPP5C			3820.2	3541.54	2230.17	4003.55	3100.01		2713.82
P53350	P53350	PLK1_HUMAN	PLK1	12582.1	12246.7	5418.79	8199.02	4448.81	3546.82	6332.92		4903.45
P53367	P53367	ARFP1_HUMAN	ARFIP1						719.026	799.173	521.08	1522.85
P53396	P53396	ACLY_HUMAN	ACLY	19990	16538.8	18963.6	14905.3	14030.1	20482.9	14620.2	7194.52	13581.4

P53582	P53582	MAP11_HUMAN	METAP1									
P53597	P53597	SUCA_HUMAN	SUCLG1		9800.06	6361.15	5355.26		3885.38	6366.7	8352.6	5975.5
P53618	P53618	COPB_HUMAN	COPB1	33496.6	34259.7	41992.5	69340.8	78235.7	74891.9	61837.8	60831.3	57251.3
P53621	P53621	COPA_HUMAN	COPA	38564.2	43112.4	34505.1	41965.8	38939.4	36502	44595.8	40382.3	43431.5
P53634	P53634	CATC_HUMAN	CTSC	9647.01	5584.49	18843.8	4793.44	2691.24	8442.88	7814.33	8746.77	9572.6
P53801	P53801	PTTG_HUMAN	PTTG1IP		902.38			856.318	1783.57	863.306		1355.06
P53985	P53985	MOT1_HUMAN	SLC16A1		2525.1	3602.21	4518.78	3355.64	3211.31	19389.1	9022.49	15378.1
P53990	P53990	IST1_HUMAN	IST1			2052.85		5175.19	4641.61	5016.25	2387.01	3855.52
P53992	P53992	SC24C_HUMAN	SEC24C	9303	7903.21	8060.29	14375	10722	16431.2	10878.3	4682.44	8324.97
P53999	P53999	TCP4_HUMAN	SUB1	43587.3	50866.6	3798.46	53255.4	46956.1	8326.93	8775.1	26611.9	7619.18
P54098	P54098	DPOG1_HUMAN	POLG	13405.7	20413		4326.29	3258.27	1724.58	1777.14	3545.15	2346.25
P54132	P54132	BLM_HUMAN	BLM		4781.27		1806.72	2474.06			1751.42	1163.83
P54136	P54136	SYRC_HUMAN	RARS1	23166.5	26820.6	23965.6	48528.8	26206.5	32558.1	18597.1	22345.5	23819.3
P54289	P54289	CA2D1_HUMAN	CACNA2D1									
P54577	P54577	SYYC_HUMAN	YARS1			4283.83	5942.98	7027.38	15752.5	6908.69	3135.15	7458.55
P54578	P54578	UBP14_HUMAN	USP14				1266.79	935.532	1593.89	1198.72		
P54619	P54619	AAKG1_HUMAN	PRKAG1					4580.44	4071.12	725.078		
P54652	P54652	HSP72_HUMAN	HSPA2	4677.55		4927.11		1739.35		1646.89	7458.32	
P54687	P54687	BCAT1_HUMAN	BCAT1				1422.7		3948.07			
P54709	P54709	AT1B3_HUMAN	ATP1B3		3398.97	6036.53	4952.81	6193.53	9786.6	28621	8377.35	21197.6
P54727	P54727	RD23B_HUMAN	RAD23B	11432.1	20503.5	4349.24	3688.12	6183.53	9827.62	5346.61	4366.76	4880.71
P54803	P54803	GALC_HUMAN	GALC									
P54819	P54819	KAD2_HUMAN	AK2	4005.81		4818.35	2545.42	2793.19	8547.84	6508.88	2211.1	5627.03
P54886	P54886	P5CS_HUMAN	ALDH18A1	9465.14	12844.9	7941.04	10385	9694.31	9272.6	20622.3	25512.7	27995.3
P54920	P54920	SNAA_HUMAN	NAPA					4487.39	3214.24	2489.91		2824.17
P55010	P55010	IF5_HUMAN	EIF5				2671.73	1964.57	2991.11	2516.2	2462.58	1785.9
P55011	P55011	S12A2_HUMAN	SLC12A2					1965.06	1560.27	17778.3	6973.71	11901.7
P55036	P55036	PSMD4_HUMAN	PSMD4			6854.71	7264.3	6566.68	10444.3	6927.4	3355.16	6963.81
P55039	P55039	DRG2_HUMAN	DRG2				4852.52			5645.85	3591.26	5064.45
P55060	P55060	XPO2_HUMAN	CSE1L	11559.2	12974.1	13201.8	36717.3	20331.3	16478.5	22237.4	37398.1	20269.1
P55061	P55061	BI1_HUMAN	TMBIM6	2315.1				1576.2	1891.81	1769.28		2027.95
P55072	P55072	TERA_HUMAN	VCP	15836	6655.35	16626.3	11358.5	20234.9	30951	24726.5	6793.01	18914.7
P55081	P55081	MFAP1_HUMAN	MFAP1				3419.2	5563.69	1801.06		2453.73	
P55084	P55084	ECHB_HUMAN	HADHB	21056.7	16487.6	16076.7	9535.33	12080.5	15062.7	24100.3	20961.6	15560.3
P55145	P55145	MANF_HUMAN	MANF	4071.15		1526.02	1812.6	2384.76	2481.54	968.157		1375.38
P55209	P55209	NP1L1_HUMAN	NAP1L1	9083.53	4847.66	9313.21	23370.2	14900.1	25268.5	11344.1	7345.47	14400.6
P55263	P55263	ADK_HUMAN	ADK					497.294	1868.8	1583.01		1053.67
P55265	P55265	DSRAD_HUMAN	ADAR	2309.6	8655.47	8746.47	1694.16	6481.68	3801.92	1812.92		1385.63
P55327	P55327	TPD52_HUMAN	TPD52							1567.28	4241.91	1964.12
P55735	P55735	SEC13_HUMAN	SEC13			7269.14					7744.06	
P55769	P55769	NH2L1_HUMAN	SNU13	2339.46	1921		2933.18	2215.26	2909.31	2484.99	2155.33	1798.58
P55786	A6NEC2;P55786	PSA_HUMAN	NPEPPS	13180.2	13701.8	13153.8	18225.6	12001.3	15728.9	8316.61	8352.62	7316.39
P55795	P55795	HNRH2_HUMAN	HNRNPH2	4191.6	2946.88	2086.35	3580.1	5273.68	3094.65	1846.52	2611.05	2495.29
P55809	P55809;Q9BYC2	SCOT1_HUMAN	OXCT1	7230.97		5524.55	3766.78	2326.97	6229.12	10779.3	7220.33	9113.8
P55884	P55884	EIF3B_HUMAN	EIF3B	8653.72	5571.11	8286.43	11511.5	9215.63	15655.2	13395.9	7210.8	12044.6
P56134	P56134	ATPK_HUMAN	ATP5MF	6818.59	7138.37	4892.9	2554.23	10875.9	4746.07	9134.3	3934.84	5275.98
P56192	P56192	SYMC_HUMAN	MARS1	15276.4	19891.9	17995.3	28832.9	36842.3	31627.6	29094.5	18496.6	21184.7
P56270	P56270	MAZ_HUMAN	MAZ					2493.04	1405.62			714.897
P56385	P56385	ATP5I_HUMAN	ATP5ME	3518.94						711.572	2328.68	
P56524	P56524	HDAC4_HUMAN	HDAC4						833.325			
P56537	P56537	IF6_HUMAN	EIF6	4794.47		7830.54	11217.1	12652.4	14308.6	12292.4	11364.3	11418.7
P56556	P56556	NDUA6_HUMAN	NDUFA6	4626.35			2166.44	2764.95	2208.33	4514.81	6065.63	5143.28
P56945	P56945	BCAR1_HUMAN	BCAR1				2728.19	3562.53	3454.09	3726.16	2092.49	3367.02
P57053	P57053;Q99879;Q5QNW6;P62807;O60814;Q99877;Q5	H2BFS_HUMAN	H2BC12L									
P57088	P57088	TMM33_HUMAN	TMEM33	8949	10136	10093.8	10273.9		4801.09	7696.51	9189.39	8925.08
P57678	P57678	GEMI4_HUMAN	GEMIN4				2674.64	3987.21	2946.81	4188.87	3532.13	4554.97
P57740	P57740	NU107_HUMAN	NUP107				1374.22	2891.1	2578.5	2839.04	801.568	1844.43
P57764	P57764	GSDMD_HUMAN	GSDMD				4993.65		3178.63	2672.04	3180.81	3192.84
P58107	P58107	EPIPL_HUMAN	EPPK1	14000.5	36723.7	12500.2		20716.3	14727.6	17433.7		19511.9
P58546	P58546	MTPN_HUMAN	MTPN				5504.37		3878.92		6766.43	3128.35
P59190	P59190	RAB15_HUMAN	RAB15				11341.4			11797.5	15291.3	15997.3
P59665; P59666	P59666;P59665	DEF1_HUMAN;DEF3_HUMAN	DEFA1B;DEFA3	19086.1	16845.1	103977	9194.3	5808.77	5182.99	5333.93	8967.15	9400.09
P59768	P59768	GBG2_HUMAN	GNG2							13283.1	4071.64	14321.6
P59998	P59998	ARPC4_HUMAN	ARPC4	8035.51	4783.23	10229.8	7302.42	5070.37	7755.85	4865.73	5522.78	6547.12
P60033	P60033	CD81_HUMAN	CD81					2708.27	2434.86	689.156		
P60174	P60174	TPIS_HUMAN	TPI1	33626.4	7721.19	45444.5	18972.7	18415.5	68535.6	33038.3	13114.3	32513.9
P60228	P60228	EIF3E_HUMAN	EIF3E	6489.45	3705.53	4865.4	9032.88	13105.3	16372.8	16356.7	5232.29	14839.6
P60468	P60468	SC61B_HUMAN	SEC61B				4006.6	2821.8	1914.12	2007.9	4391.64	2210.29
P60660	P60660	MYL6_HUMAN	MYL6	79726.4	56836.5	58289.3	268860	83569.2	71523.5	71503.2	196753	61573.4
P60709; P63261	P60709;P63261	ACTB_HUMAN;ACTG_HUMAN	ACTB;ACTG1	##########	893966	998383	1.30666E+06	2.23496E+06	1.29191E+06	1.66209E+06	954849	1.14118E+06
P60842	P60842	IF4A1_HUMAN	EIF4A1	40758.1	27858.1	49110.3	69413.4	79560	125883	114274	65316.3	91482.5
P60866	P60866	RS20_HUMAN	RPS20	114872	108579	110203	42615.3	115953	74983.5	72839.9	49376.2	66094
P60891	P21108;P60891;P11908	PRPS1_HUMAN	PRPS1	7587.86		7245.08	7825.3	5786.23	7838.32	9099.59	7757.3	9377.56
P60900	P60900	PSA6_HUMAN	PSMA6	4563.17	5594.69	11691.5	4887.14	11541.9	20454.8	13816.7	1866.6	9994.3
P60903	P60903	S10AA_HUMAN	S100A10	9629.05	9965.02	7567.4	7459.81	7891.42	11287.9	7436.63	6087.5	6352.72
P60953	P60953	CDC42_HUMAN	CDC42				6185.76	8180.82	4760.84	6163.88	6261.53	6140.38
P60981	P60981	DEST_HUMAN	DSTN	3431.95		4821.75	2488.55	3424.88	3531.91	2910.98	5859.37	2609.56
P61006	P61006	RAB8A_HUMAN	RAB8A	3741.86		6733.89		5268.15	6066.06	8017.26	2142.4	8404.92
P61011	P61011	SRP54_HUMAN	SRP54				2111.11		2664.71	2017.74	1815.85	2515.27
P61019	P61019;Q8WUD1	RAB2A_HUMAN	RAB2A	5616.99		7852.34	5168.89	10733.5	5162.47	6022.5	4331.7	7204.01
P61020	P61020	RAB5B_HUMAN	RAB5B					1542.11	1281.96	1306.6		1866.05
P61026	P61026	RAB10_HUMAN	RAB10			3451.15		8858.82	5505.87	11350.8	4908.81	11880.7
P61077	P62837;P61077	UB2D3_HUMAN	UBE2D3	5368.48		5871.93	6066.21	1311.92	2003.75		3694.91	
P61088	P61088	UBE2N_HUMAN	UBE2N	6345.6		13775.4	5138.47	12258.8	8726.57	3511.42	5221.56	4880.19
P61106	P61106	RAB14_HUMAN	RAB14	15874.7	8237.15	10695.3	5884.83	6971.93	6851.99	5748.11	5773.09	5233.47
P61158	P61158	ARP3_HUMAN	ACTR3	5445.82	4990.22	2843.65	5250.74	6777.94	10894.2	4147.24	6675.37	3201.05
P61160	P61160	ARP2_HUMAN	ACTR2			28461.2	6590.18	5686.25	5400.04	5512.49	7556.49	6861.27
P61163	P61163;P42025	ACTZ_HUMAN	ACTR1A	4420.33	4958.88	6322.96	2855.86	9787.69	5812.48	4691.21	2657.58	4926.08
P61201	P61201	CSN2_HUMAN	COPS2	6129.1	3371.64	2152.98	3625.36	3959.47	5648.2	4815.1	1948.11	4110.06
P61204; P84077	P61204;P84077	ARF1_HUMAN;ARF3_HUMAN	ARF1;ARF3	4533.05	3016.99	3200.88	3453.61	1122.61	3317.03	2195.14	2115.11	1832.25
P61221	P61221	ABCE1_HUMAN	ABCE1		5429.87	4065.15	4420.23	4672.17	6550.78	3853.7	2731.99	4581.42
P61224	P61224;A6NIZ1	RAP1B_HUMAN	RAP1B				3660.08	6134.11	6614.11	7381.04	3456.23	7564.99
P61225	P61225	RAP2B_HUMAN	RAP2B				6184.49	16752.1	13317.8	28768.1	4179.27	45913.5
P61247	P61247	RS3A_HUMAN	RPS3A	70876	106926	76876.9	104227	149907	123271	113389	85240.1	92027.9
P61254	P61254;Q9UNX3	RL26_HUMAN	RPL26	16783.5	25738.7	21276.6	28227.9	41014.3	36275.3	27549	31217.2	28897.8
P61289	P61289	PSME3_HUMAN	PSME3			3347.54	5280.61	3909.43	5920.48	5138.19	4686.95	5512.93
P61313	P61313	RL15_HUMAN	RPL15	12523.2	15186.5	14057.3	18935.1	33886.1	34623.6	22644.4	15823.8	24304.2
P61353	P61353	RL27_HUMAN	RPL27	22961.9	36752.6	20353.2	40379.4	61962	50628.4	55421.6	48837.9	52407.2
P61421	P61421	VA0D1_HUMAN	ATP6V0D1				14768		1920.27	2992.56	13080	2485.15
P61457	P61457	PHS_HUMAN	PCBD1			5337.95			2062.11	12758	9448.82	11847.8
P61513	P61513;A6NKH3	RL37A_HUMAN	RPL37A	19668.8	10202.5	10155.8	40209.3	13713.3	16049.1	21522.5	48312.6	13168.4
P61586	P08134;P62745;P61586	RHOA_HUMAN	RHOA	8979.21	4800.04	8185.82	6258.42	8683.97	11866.9	20561.6	9774.47	19820.2
P61604	P61604	CH10_HUMAN	HSPE1	29398.8	10045.5	18473.4	18402.7	15329.3	35403.1	25007.3	18567.8	24060.4
P61619	Q9H9S3;P61619	S61A1_HUMAN	SEC61A1	9940.43	10986.7	11293.3	9632.22	4347.74	6524.61	4094.78	8188.31	4148.32
P61626	P61626	LYSC_HUMAN	LYZ	39088.8	26462.5	60220.6	29006.9	8769.58	50541.7	28160.6	48435.2	20275.4
P61758	P61758	PFD3_HUMAN	VBP1							1659.47		
P61769	P61769	B2MG_HUMAN	B2M	3806.28		3411.25	2785.05	3508.66	2221.7	4296.71	2341.99	3687.8
P61803	P61803	DAD1_HUMAN	DAD1				3929.42			2159.22	6005.51	
P61923	P61923	COPZ1_HUMAN	COPZ1			895.687		2609.61	2260.24	640.658		
P61956	Q6EEV6;P61956;P55854	SUMO2_HUMAN	SUMO2				2500.86		4486.14	3059.54	3426.84	3233.07
P61960	P61960	UFM1_HUMAN	UFM1	4726.24		5719.98	3178.74		2804.41	2730.85	4438.8	3966.35
P61962	P61962	DCAF7_HUMAN	DCAF7				1293.57	2987.08	2614.12	2685.54	2873.5	3802.75
P61966	P61966	AP1S1_HUMAN	AP1S1			3703.74	2613.42				2789.41	
P61978	P61978	HNRPK_HUMAN	HNRNPK	26206.7	12835.7	21945.9	31986.7	29560.4	32944.3	21252.8	18371.4	16728.5
P61981	P61981	1433G_HUMAN	YWHAG	7267.96		9198.72	4954.35	3168.56	8353.73	7059.35	5785.48	8212.64
P62070	P62070	RRAS2_HUMAN	RRAS2				3938.66	12137.8	19622.2	13550.9	3665.42	11951.2
P62081	P62081	RS7_HUMAN	RPS7	12363.4	11975	15873.6	12020.9	49890.3	26879.4	24074.9	17469.1	26954
P62136	P36873;P62140;P62136	PP1A_HUMAN	PPP1CA	8680.98	8162.02	13980.6	8541.46	23564.9	23078.3	8499.98	5188.54	8330.59
P62140	P62140	PP1B_HUMAN	PPP1CB	5258.65		23542.1	6187.47	30895.8	20324.8	4912.66	1173.85	
P62166	P62166	NCS1_HUMAN	NCS1						3233.45	6382.16		7284.71
P62191	P62191	PRS4_HUMAN	PSMC1	25025.5	11158.2	21126.5	33523.6	22110.3	18339.3	18674.6	16772.4	16456.3
P62195	P62195	PRS8_HUMAN	PSMC5	13363.2	13868.4	12678.5	15575.7	14832.3	14128.6	16363.5	16493.5	20132.4
P62241	P62241	RS8_HUMAN	RPS8	32371.6	31024.3	27457.9	38371.9	38523.1	32900.2	36156.7	39586.7	33985
P62244	P62244	RS15A_HUMAN	RPS15A	62153.3	86014.1	48316.6	55416.8	78523.8	47193.1	50041.1	82985.8	55494.1
P62249	P62249	RS16_HUMAN	RPS16	214642	235785	205448	179824	336556	186204	189660	187199	195045
P62256	P62256	UBE2H_HUMAN	UBE2H				2048.46	4182.02	7893.93			
P62258	P62258	1433E_HUMAN	YWHAE	12634.5	9415.74	11748.6	8003.56	9143.07	22227.2	18217.3	7392.08	19784.9
P62263	P62263	RS14_HUMAN	RPS14	44022.9	56539.7	50738	47172	36830.7	25266.7	41395.3	51941.4	40660
P62266	P62266	RS23_HUMAN	RPS23	52658	41392.4	42945.9	46163.5	20693.5	9856.28	19003.9	43729.6	23798.1
P62269	P62269	RS18_HUMAN	RPS18	140231	225104	129808	132547	234692	136753	125391	132756	142433
P62273	P62273	RS29_HUMAN	RPS29	12365.6	14133	12792.9	8250.43	6686.71	5092.01	6918.2	10945.9	6887.27
P62277	P62277	RS13_HUMAN	RPS13	54848.1	64544.5	60415.7	66070.9	126230	62527.2	58403.4	60793.9	65468.8
P62280	P62280	RS11_HUMAN	RPS11	99297.5	108718	171487	80609.1	135239	98514.5	123709	84921.5	168689
P62304	P62304	RUXE_HUMAN	SNRPE	10413.4	10721.5	4903.98	4932.15	26340.3	9887.68	20391.1	14718.1	17448.1
P62310	P62310	LSM3_HUMAN	LSM 3:00				1496.55				5082.76	
P62314	P62314	SMD1_HUMAN	SNRPD1				4278	7087.48	4679.4	5408.04	6098.95	5404.5
P62316	P62316	SMD2_HUMAN	SNRPD2				4324.86	9180.72	7310.5	2201.88	8904.59	2860.85
P62318	P62318	SMD3_HUMAN	SNRPD3		25679		2350.75	30816.7	10208	7317.37	13607.6	5467.69
P62328	P62328	TYB4_HUMAN	TMSB4X	7784.89		4659.04	8113.06	1564.2	5151.1	5932.42	12118.4	5589.62
P62330	P62330	ARF6_HUMAN	ARF6	8147.54		11804.3	7875.44			4896.96	5544.81	7178.77
P62333	P62333	PRS10_HUMAN	PSMC6	15411.5	16536.6	15026.6	19888.3	16944.5	17994	12212.8	13832.2	11441.5
P62424	P62424	RL7A_HUMAN	RPL7A	26964.3	34302.4	28635.9	34262.6	55144.4	41698.7	40939.8	34862	35211.7
P62495	P62495	ERF1_HUMAN	ETF1	8771.07		8656.43	18867.5	5919.38	5316.58	7394.29	14954.7	6554.81
P62633	P62633	CNBP_HUMAN	CNBP			2635.9	716.222	1724.22	2794.71	682.97	2054.71	1463.66
P62701	P62701;P22090;Q8TD47	RS4X_HUMAN	RPS4X	65853.1	63416.6	63232.8	68802.3	78752	55879.9	56921.8	60681.6	62244.5
P62745	P62745	RHOB_HUMAN	RHOB						653.147	4886.29		4867.81
P62750	P62750	RL23A_HUMAN	RPL23A	40201.5	43160.4	38851.2	46780.8	89401.6	55358.9	56631	52272.3	51725.7
P62753	P62753	RS6_HUMAN	RPS6	35736.2	39670.7	31026.9	33318.8	46956	28913.2	30980.3	30075.2	30819.6
P62760	P62760	VISL1_HUMAN	VSNL1							5018.63	4589.8	4647.98
P62805	P62805	H4_HUMAN	H4C16	803378	594697	346333	366059	523156	185738	109227	179912	77832.6
P62820	P62820	RAB1A_HUMAN	RAB1A	4450.87	4424.26	7191.63	3346.71	8906.02	6658.42	6781.52	2756.77	4735.43
P62826	P62826	RAN_HUMAN	RAN	5496.31		11305.5	3963.75	9423.22	12716.3	5339.86	2465.92	4321.3
P62829	P62829	RL23_HUMAN	RPL23	39890.8	70540.8	66769.4	40775.8	71788.7	44415.5	38575.3	53658.4	45212.5
P62834	P62834;P61224	RAP1A_HUMAN	RAP1A	3102.05			5705.27	15124.8	7832.02	8863.13	5218.94	7926.78
P62841	P62841	RS15_HUMAN	RPS15	9728.87	9387.52	12027.2	7345.23	32788.8	20898.8	11594	7645.91	15661.1
P62847	P62847	RS24_HUMAN	RPS24	11419.7	12663.9	15699.9	10112.7	27324	16336.4	15217.4	13195.3	20412
P62851	P62851	RS25_HUMAN	RPS25	73029.7	95761.4	46200.9	57831.9	63323.4	46885.2	48587.1	52729.2	53355.9
P62854	Q5JNZ5;P62854	RS26_HUMAN	RPS26	16993.1	18847.3	13119	11432.3	31313	17058.9	17967.8	24324.6	20018.3
P62857	P62857	RS28_HUMAN	RPS28	22130	27125	19967.9	17875.5	31099.8	22776.8	10509.9	7198.66	13614.9
P62861	P62861	RS30_HUMAN	FAU	53409.2	58195.3	34051.6	31478.2	61359.4	17784.4	37503.7	49019	31638.1
P62873	P62873;P62879	GBB1_HUMAN	GNB1	7897.65	13326.3	11257.5	40403.3	34118.6	42466.3	135321	40146.7	138273
P62879	P62879	GBB2_HUMAN	GNB2			2561.2	9625.87	7429.28	9498.61	22380.1	5670.41	24706.6
P62888	P62888	RL30_HUMAN	RPL30	36491	36247.9	26279.6	36187.3	46666.5	44452.9	35029.1	48382.1	35412.6
P62891; Q59GN2	P62891;Q59GN2	R39L5_HUMAN;RL39_HUMAN	RPL39;RPL39P5	13699.7	16985.1	15135.8	7746.2	9932.62	6786.68	5964.82	6966.87	6728.08
P62899	P62899	RL31_HUMAN	RPL31	23877.7	24645.2	26735.6	23508.6	45638.5	40429.2	22756.4	35223.9	26045.6
P62906	P62906	RL10A_HUMAN	RPL10A	27169	46891.8	32163.8	33067.1	50323.1	49037.4	48327.1	29100.4	44407.6
P62910	P62910	RL32_HUMAN	RPL32	8639.1	7762.46	8632.58	12084.2	13565.5	13829.4	10477.3	9948.01	8735.55
P62913	P62913	RL11_HUMAN	RPL11	19879.6	27442.6	31397.7	37883.3	26848.8	35864.1	40435.1	35895.5	35698.9
P62917	P62917	RL8_HUMAN	RPL8	21804.2	26309.3	19882.7	33800.7	48497.2	33302.9	35798	29982.9	31862.6
P62937	P62937	PPIA_HUMAN	PPIA	56192.9	19067.3	55131.5	61555.2	40933.7	68813.7	36718.4	21289.9	31287.9
P62942	P62942	FKB1A_HUMAN	FKBP1A	5358.33		2996.75	2669.77	3411.63	7207.97	3762.13	1562.62	2296.94
P62979	P62979	RS27A_HUMAN	RPS27A				14617.3	9228.02	11509.6	11037.2	20682.5	10127.8
P62987	P0CG47;P0CG48;P62979;P62987	RL40_HUMAN	UBA52	181848	177581	171488	179486	137363	150136	145412	156041	166285
P62995	P62995	TRA2B_HUMAN	TRA2B		9686.43			9136.52	3483.35	4248.56		4864.92
P63000	P15153;P60763;P63000	RAC1_HUMAN	RAC1			8607.16	8912.93	7098.66	10037.5	11890.7	4629.51	12950.8
P63010	Q10567;P63010	AP2B1_HUMAN	AP2B1	5233.93	6970.2	3762.02	7322.95	12126	7818.88	6436.02	4306.56	4557.36
P63096	P63096	GNAI1_HUMAN	GNAI1	5838.3			11495.4	5576.75	8027.63	78470.8	23581.6	86400.8
P63104	P63104	1433Z_HUMAN	YWHAZ	21565.5	13182.1	27903.4	26663.9	42217.1	124137	44009.9	17691.6	53602.3
P63151	P63151	2ABA_HUMAN	PPP2R2A				2721.37	2490.21	2815.79	1774.69	1498.09	1783.58
P63167	P63167;Q96FJ2	DYL1_HUMAN	DYNLL1	6166.22	10838	7144.47	5848.43	7499.14	6968.91	10704	6866.07	7814.95
P63172	P63172	DYLT1_HUMAN	DYNLT1	3249.31	1231.26	950.117	2839.98	1637.45	1611.59	3404.47	4036.6	1595.54
P63173	P63173	RL38_HUMAN	RPL38	46451.3	52438.9	41125.3	46624.6	56133.6	47996.2	46571.8	34433.9	27179.8
P63208	P63208	SKP1_HUMAN	SKP1	10140	22775.7		6105.06	7184.34	8453.3	5351.01	5061.66	5083.88
P63218	P63218	GBG5_HUMAN	GNG5							5140.99		5554.49
P63220	P63220	RS21_HUMAN	RPS21	3684.64	4730.64	3178.11	3704.83	28111.8	13496.9	10364.8	5312.89	9148.67
P63241	Q9GZV4;P63241;Q6IS14	IF5A1_HUMAN	EIF5A	3955.26	2082.91	4264.05	5324.99	3662.09	13618.4	4182.84	2154.82	4356.3
P63244	P63244	RACK1_HUMAN	RACK1	14965.5	18068.9	17747.3	16088	21846.3	21086.9	12522.3	14037.6	13213.5
P63267	P62736;P63267;P68032	ACTH_HUMAN	ACTG2	8984.44	315229	162987	1214.82	218942	66218.7	479370		
P63279	P63279	UBC9_HUMAN	UBE2I			3847.94	3508.14		3056.06	2772.36	1932.19	2347.68
P63313	P63313	TYB10_HUMAN	TMSB10	30668.9	13438.3	22236.3	16419.5	3535.89	12442.8	12825.1	18258.8	16087.8
P63316	P63316	TNNC1_HUMAN	TNNC1									
P67775	P67775;P62714	PP2AA_HUMAN	PPP2CA	5955.32	6903.67	6697.37	11180.9	13178.5	8298.01	7516.2	8656.27	12238
P67809	P16989;P67809;Q9Y2T7	YBOX1_HUMAN	YBX1	11530.3	15589.7	8827.54	20255.2	28510.3	16803.2	17710.8	33935.3	17396
P67812	P67812	SC11A_HUMAN	SEC11A					3975.32	1022.21	1593.86		555.996
P67936	P67936	TPM4_HUMAN	TPM4	24763.1	23093.1	23502.1	34301.3	90749.5	55935.4	30208.9	18289.6	27228.6
P68036	P68036	UB2L3_HUMAN	UBE2L3			2094.69	1809.38	4184.56	7352	1226.48		1506.38
P68104	P68104;Q5VTE0	EF1A1_HUMAN	EEF1A1	571629	428771	474635	1.02109E+06	558046	551172	1.0136E+06	1.09508E+06	881714
P68133	P62736;P68133;P63267;P68032	ACTS_HUMAN	ACTA1	57737.2	102610	69183.1	61855.3	547881	101885	86528.1	30143.9	73517.8
P68366	P68366	TBA4A_HUMAN	TUBA4A	171847	208686	152017	212216	114056	108067	285657	316881	284619
P68371	P04350;P68371	TBB4B_HUMAN	TUBB4B	324668	230455	267229	373126	183631	137338	337953	577719	378445
P68400; Q8NEV1	Q8NEV1;P68400	CSK21_HUMAN;CSK23_HUMAN	CSNK2A1;CSNK2A3	11137.4	1593.75	2474.04	2875.02	13631.2	8688.76	7043.4	3701.12	2891.5
P68402	P68402	PA1B2_HUMAN	PAFAH1B2	6510.31	2067.9	4733.67	12049.6	22263.3	36852.3	15411.7	7816.74	7228.5
P68871	P68871;P02042	HBB_HUMAN	HBB	20419.1	11643.3	17087.4	8461.41	13797.6	3589.51	10789.7	8932.71	12626.6
P69849; Q15155; Q5JPE7	Q5JPE7;P69849;Q15155	NOMO1_HUMAN;NOMO2_HUMAN;NOMO3_	NOMO1;NOMO2;NOMO3	4767.57	4729.14	6616.99	5684.05	17317.6	12483.3	10628.8	5016.7	7530.9
P69905	P69905	HBA_HUMAN	HBA2	22688	18319.1	24710.8	5018.88	3372.57	3000.99	7730.99	6182.1	9347.58
P78310	P78310	CXAR_HUMAN	CXADR				5835.35	3432.48	3833.93	7461.15		5254.28
P78318	P78318	IGBP1_HUMAN	IGBP1	6002.37	5416.84	5895.36	5422.45	3999.5	7734.22	7747.51	7559.72	8432.39
P78344	P78344	IF4G2_HUMAN	EIF4G2	13163.8	8153.82	15717.9	12597.6	9272.22	12453.6	11433.1	10777.2	10755.3
P78347	P78347	GTF2I_HUMAN	GTF2I				3563.15	2068.7	2477.91	7097.26	5059	6651.31
P78358	P78358	CTG1B_HUMAN	CTAG1B				21947.3	43602.7	17346	950.412	3176.55	
P78368	P78368	KC1G2_HUMAN	CSNK1G2					4874.31	5558.39	17540.2	2044.84	14365
P78371	P78371	TCPB_HUMAN	CCT2	11169.8	7236.72	10809.7	18418.8	27544.3	34162	32285.5	11450.4	29631.9
P78385	P78385	KRT83_HUMAN	KRT83			1307.41						750.464
P78386	P78386	KRT85_HUMAN	KRT85	66212.5	83772.4	78527.5	44530.4	4994.35	23258.7	42301.3	54089.8	62739.6
P78395	P78395	PRAME_HUMAN	PRAME				1895.45	2683.56	2451.28			
P78406	P78406	RAE1L_HUMAN	RAE1				1924.17	3528.66	2259.83	2745.02	1052.35	2995.41
P78417	P78417	GSTO1_HUMAN	GSTO1	29537.1	25815.2	39767.1	20825.3	13649	52095.3	15576.5	20129	16098.5
P78524	P78524	DEN2B_HUMAN	DENND2B	2105.75	1235.47	554.142	1608.08	3640.67	2146.12			
P78527	P78527	PRKDC_HUMAN	PRKDC	25258.1	28117.9	16429	28905.5	28645.7	22219.3	26636.7	34161	30702.4
P78549	P78549	NTH_HUMAN	NTHL1		2790.24	2136.95		3324.86				
P78559	P78559	MAP1A_HUMAN	MAP1A		3241.29		1735.99	2199.1	1819.53			
P80188	P80188	NGAL_HUMAN	LCN2	4912.51							4826.31	
P80723	P80723	BASP1_HUMAN	BASP1	19586.6	13300.5	13477.6	59134.1	59858.5	66826.8		2298.67	851.946
P81605	P81605	DCD_HUMAN	DCD	29687.4	24727.1	41116.7	23255.1	6127.14	8660.02	9968.26	27544.5	12124.9
P82650	P82650	RT22_HUMAN	MRPS22					2969.54	2113.33	2660.2		2213.6
P82909	P82909	KGD4_HUMAN	MRPS36			2430.81	1253.9	1482.17	1313.31	5023.56	1555.24	4782.93
P82912	P82912	RT11_HUMAN	MRPS11	4284.07	6623.41	6648.22	2126.69		1318.74	3077.96		4101.99
P82930	P82930	RT34_HUMAN	MRPS34	2945.1		4151.92	2606.4	3570.71	3806.24	5895.56	4248.73	7576.35
P82933	P82933	RT09_HUMAN	MRPS9			1744.88		4285.26	1869.75	4658.99	2049.02	5675.16
P82970	P82970	HMGN5_HUMAN	HMGN5						885.413	4502.95		3558.19
P82979	P82979	SARNP_HUMAN	SARNP	5081.72		9200.29	5504.48	4899.89	5853.15	5776.43	6627.39	6372.06
P83111	P83111	LACTB_HUMAN	LACTB					3036.45	653.74			
P83436	P83436	COG7_HUMAN	COG7	1308.19	1379.03		1985.25	1528.44	1282.18	1626.15	1137.39	1207.82
P83731	P83731	RL24_HUMAN	RPL24	33987.4	44003.5	31941.1	34771.6	47458.3	26367	24419.2	39481.1	27797.3
P83881	P83881;Q969Q0	RL36A_HUMAN	RPL36A	12768.9		12543.5	11442.4	5736.36	6281	7532.92	15827.5	8346.5
P84085	P84085	ARF5_HUMAN	ARF5									
P84090	P84090	ERH_HUMAN	ERH					4486.48	2281.25			
P84095	P84095	RHOG_HUMAN	RHOG		2906.27		6841.52	5237.69	2983.58	6296.76	7497.24	6572.46
P84098	P84098	RL19_HUMAN	RPL19	10785.3	10360.6	8872.58	12580.4	16393.2	7871.03	12056.1	12284.8	11297.2
P84103	P84103	SRSF3_HUMAN	SRSF3	16723	26501.8	13992.9	9251.76	29224.1	16179.3	12721.3	11654.3	14547.2
P84243	P84243	H33_HUMAN	H3-3B	98473	109880	56036.6	93431.8	114385	52932.6	21248.6	43176.1	18776
P85037	P85037	FOXK1_HUMAN	FOXK1	10683.5	13558.1	6797.32	7551.81	18709.7	8707.95	8387.32	6123.82	7639.3
P98164	P98164	LRP2_HUMAN	LRP2	32653.9	35561	14491.8	30126.8	3361.08	2481.94	7213.43	28473.3	6273.92
P98171	P98171	RHG04_HUMAN	ARHGAP4							7398.03	8660.55	11238.7
P98172	P98172	EFNB1_HUMAN	EFNB1							2772.7	945.967	3015.51
P98179	P98179	RBM3_HUMAN	RBM3					1880.73	1112.95	1355.85	2104.93	1072.93
P99999	P99999	CYC_HUMAN	CYCS	16218.1		14320.1	7420.24	8317.93	31658.1	10784.2	5771.41	10420.6
Q 00013	Q00013	EM55_HUMAN	MPP1	7774.07			5043.86					
Q 00059	Q00059	TFAM_HUMAN	TFAM	77172.3	104652	39262	80338.5	105481	75512	21951	29046.4	20379
Q 00169	Q00169	PIPNA_HUMAN	PITPNA		980.909	3569.35		5840.4	6634.09	2986.83	1409.84	2900.22
Q 00325	Q00325	MPCP_HUMAN	SLC25A3	6795.97	11824.1	8304.19	3994.31	4646.05	5595.21	10147.8	7329.22	8556.38
Q 00341	Q00341	VIGLN_HUMAN	HDLBP	4247.64			5760.95	6426.76	5932.73	4344.67	4000.69	4638.76
Q 00534	Q00534	CDK6_HUMAN	CDK6				3324.58	4404.12	4909.54	3200.08	3517.77	5290.82
Q 00537	Q07002;Q00537	CDK17_HUMAN	CDK17		1268.23	840.86	854.922	1549.04	1355.6			
Q 00577	Q00577	PURA_HUMAN	PURA	22153.4	36060.2		16310.3	15485.7	6727.64	5656.58	13522	3649.04
Q 00587	Q00587	BORG5_HUMAN	CDC42EP1				5630.12	2779.96	2063.17	1290.98	1762.46	4211.22
Q 00610	Q00610	CLH1_HUMAN	CLTC	80997.3	93208.7	68077.6	110409	106073	106751	70900.9	44339.4	69005.8
Q 00688	Q00688	FKBP3_HUMAN	FKBP3				1490.26	2475.05	7385.8	3243.49		3338.5
Q 00796	Q00796	DHSO_HUMAN	SORD			3240.27			1325.52	3581.15	4919.71	3644.98
Q 00839	Q00839	HNRPU_HUMAN	HNRNPU	22115.2	15931.4	13212.3	12134.2	15936.4	13407.9	13843.7	14891.2	16107
Q 00872	Q00872	MYPC1_HUMAN	MYBPC1	3765.96		3330.45						
Q 01081	Q01081	U2AF1_HUMAN	U2AF1	5824.14	26389	2226.92	7611.33	84065.5	15288.1	10442.1	11215.3	6273.35
Q 01082	Q01082	SPTB2_HUMAN	SPTBN1	6444.82	7269.16	5989.08	7437.86	9349.37	10115	7896.07	3632.19	7208.68
Q 01085	Q01085;P31483	TIAR_HUMAN	TIAL1		1263.36	1238.22	556.523	12266.6	4073.79	1855.86		883.539
Q 01105	P0DME0;Q01105	SET_HUMAN	SET	8881.87	112751	9242.54	12383.3	7478.33	20588.6	14156.6	11970.9	15806.4
Q 01130	Q01130	SRSF2_HUMAN	SRSF2					15157.8	5035.55	7870.8		6812.55
Q 01167	Q01167	FOXK2_HUMAN	FOXK2					2428.68		4290.86	4342.57	
Q 01196	Q01196	RUNX1_HUMAN	RUNX1			1695.19	2370.76	6528.98	2132.75	3686.85	3244.85	4187.11
Q 01433	Q01433	AMPD2_HUMAN	AMPD2				403.44	513.631	630.802	509.358		479.418
Q 01469	Q01469	FABP5_HUMAN	FABP5	23595.2	10943.6	30982.8	8499.81	6426.74	7653.2	8798.73	12788.5	10842.5
Q 01518	Q01518	CAP1_HUMAN	CAP1	19568.9	11439.7	22108.5	21690.8	17864.4	35938.6	13221.5	5543.81	10659.1
Q 01546	Q01546	K22O_HUMAN	KRT76				1038.16					
Q 01628	Q01628	IFM3_HUMAN	IFITM3	2755.57	3484.5	2354.6	1021.87	5561.4	4822.93			688.406
Q 01650	Q01650	LAT1_HUMAN	SLC7A5	3701.26		3462.92	6219.39	5456.32	5418.35	6195.26	7641.74	11372.5
Q 01658	Q01658	NC2B_HUMAN	DR1	15567.5	12092.6	6916.56	11071.7	19080.4	16943.8	4255.05		
Q 01664	Q01664	TFAP4_HUMAN	TFAP4		2729.24							
Q 01780	Q01780	EXOSX_HUMAN	EXOSC10		2167.52			1735.9	1515.59	1694.86		777.45

Q07021	Q07021	C1QBP_HUMAN	C1QBP			2713.15	4097.46	2539.29	5297.04	4251.67	4232.42	3686.28
Q07065	Q07065	CKAP4_HUMAN	CKAP4	20422.6	14887.5	18296.6	5007.24	9945.38	8665.35	4603.53	1777.63	5023.59
Q07157	Q07157	ZO1_HUMAN	TJP1	8136.8	7827.84	3425.99	9441.64	20957.7	15264	10755.4	8294.99	11668.5
Q07352	Q07352	TISB_HUMAN	ZFP36L1	5121.32	4826.6	3759.41		2391.52		2573.79	1171.71	2325.84
Q07666	Q07666;Q5VWX1	KHDR1_HUMAN	KHDRBS1	16046.5	17690.1	15781.2	13004.8	51840.9	17957.8	19459.9	13922.9	21111.1
Q07812	Q07812	BAX_HUMAN	BAX	6511.01				5556.49	6267.02	7482.32		3670.83
Q07820	Q07820	MCL1_HUMAN	MCL1				3173.78				698.92	545.306
Q07864	Q07864	DPOE1_HUMAN	POLE						1395.19	2378.62	2487.68	2214.17
Q07866	Q07866	KLC1_HUMAN	KLC1		4845.67	3730.86	6097.15	4640.46	6121.19	4246.58	2706.12	2835.16
Q07954	Q07954	LRP1_HUMAN	LRP1	4943.67		4508.47	3211.52	3025.09	3509.25		1636.93	
Q07955	Q07955	SRSF1_HUMAN	SRSF1	27251.3	40738.4	26816.1	13618.3	39511.1	19602.7	12913.4	19397.4	12797.8
Q07960	Q07960	RHG01_HUMAN	ARHGAP1	7205.24		7688.67	6124.27	4078.65	6114.93	5465.28	4214.15	5492.9
Q07973	Q07973	CP24A_HUMAN	CYP24A1						6771.75			
Q08043	Q08043	ACTN3_HUMAN	ACTN3	5098.76			5767.36	5903.73		6104.1		9485.94
Q08188	Q08188	TGM3_HUMAN	TGM3	6098.1	13003.5	27457.1	12191.9	4578.73	7596.21	9844.71	11065.8	16966.2
Q08209	P16298;P48454;Q08209	PP2BA_HUMAN	PPP3CA			4480.91	2849.3		2324.41	2929.75		1933.58
Q08211	Q08211	DHX9_HUMAN	DHX9	7200.4	10428.3	6083.66	8334.46	24852.6	11356.7	9431.86	5621.46	8426.76
Q08257	Q08257	QOR_HUMAN	CRYZ	6325.4		4558.84	6511.61	4668.84	11169.3	5303.18	4128.91	6728.47
Q08378	Q08378	GOGA3_HUMAN	GOLGA3	1114.35	1432.39		1436.78	2942.3	2506.16	1365.95		1318.45
Q08379	Q08379	GOGA2_HUMAN	GOLGA2					3849.99	2153.23	3505.33	1372.46	3266.91
Q08380	Q08380	LG3BP_HUMAN	LGALS3BP	14288	15535.7	14844.2	24757.6	17759.1	11639.3	17076.9	32306.6	21354.8
Q08554	Q08554	DSC1_HUMAN	DSC1	25452.1	21191.4	33034.5	18283	8047.59	10460.1	9889.68	19895.9	17195.2
Q08752	Q08752	PPID_HUMAN	PPID				1638.11	3281.24	1977.22	2424.54	1205.34	2422.82
Q08945	Q08945	SSRP1_HUMAN	SSRP1	12021.4	16010	7755.43	10652.3	12516.5	9803.98	10088.5	11903.6	9004.7
Q08AD1	Q08AD1	CAMP2_HUMAN	CAMSAP2			808.759	1175.85	2186.77	2325.01	415.411		417.652
Q08AE8	Q08AE8	SPIR1_HUMAN	SPIRE1					4235.76	2552.89			
Q08AF3	Q08AF3	SLFN5_HUMAN	SLFN5	4554.62	2767.69	4043.41	2536.14	4995.43	2447.93			
Q08AM6	Q08AM6	VAC14_HUMAN	VAC14					7069.84	1526.98	3166.07	1849.26	2711.32
Q08J23	Q08J23	NSUN2_HUMAN	NSUN2	7588.14	3796.62	6205.39	7827.39	10289.5	9453.96	9338.31	6582.12	8950.86
Q09028	Q09028	RBBP4_HUMAN	RBBP4	3947.34			4200.97			1940.49	5043.44	5464.02
Q09161	Q09161	NCBP1_HUMAN	NCBP1	4621.11	2621.92		3913.69	3549.65	2658.4	2626.63	1612.61	2979.39
Q09328	Q09328	MGT5A_HUMAN	MGAT5							1982.21	1713.9	1500.32
Q09666	Q09666	AHNK_HUMAN	AHNAK	25574.8	16547.3	22045.1	19922.9	18889.4	23653.9	15864.6	12013.8	14933
Q0D2I5	Q0D2I5	IFFO1_HUMAN	IFFO1	750.638	1827.24			2686.1	2144.78			
Q0VDF9	Q0VDF9	HSP7E_HUMAN	HSPA14	2539.41		2200.5	3890.42	2364.86	2021.54	5765.44	4680	6100.27
Q0VF96	Q0VF96	CGNL1_HUMAN	CGNL1					9876.5	7120.89	16699		
Q0ZGT2	Q0ZGT2	NEXN_HUMAN	NEXN	4977.99	10608.9	4612.84	5968.42	43110.6	34708.2	11848.5	5147.77	
Q10567	Q10567	AP1B1_HUMAN	AP1B1						641.894			
Q10570	Q10570	CPSF1_HUMAN	CPSF1		2047.18		1154.81	2761.05	845.234	1652.65		1926.51
Q10588	Q10588	BST1_HUMAN	BST1				4896.79	3892.66	4594.41			
Q10589	Q10589	BST2_HUMAN	BST2						2746.12	1302.87		
Q10713	Q10713	MPPA_HUMAN	PMPCA	4508.48		5456.86	4055.2	4920.82	4761.85	1889.03		1765.42
Q12765	Q12765	SCRN1_HUMAN	SCRN1	3870.27			1287.84	1597.15	4123.05	1130.76		
Q12768	Q12768	WASC5_HUMAN	WASHC5	4001.08	4046.94	3014.17	3597.98	2516.09	3329	3635.83	2788.38	2643.25
Q12769	Q12769	NU160_HUMAN	NUP160				788.363	1247.06	1109.29	1868.88	991.89	2791.04
Q12774	Q12774;A0A2R8YFR7	ARHG5_HUMAN	ARHGEF5							3092.39	2487.23	3223.84
Q12792	Q12792	TWF1_HUMAN	TWF1	7748.16	7632.85		6075.17	5725.71	4553.66	2713.22	3872.78	2249.97
Q12797	Q12797	ASPH_HUMAN	ASPH	10901.2	11957.3	10892.1	4051.85	8391.47	4985.12	5275.14	5658.89	4695.98
Q12800	Q12800;Q9NZI7	TFCP2_HUMAN	TFCP2					3822.07	1912.87	2688.88	1787.37	3073.77
Q12824	Q12824	SNF5_HUMAN	SMARCB1							290.585		368.111
Q12834	Q12834	CDC20_HUMAN	CDC20				2640.79	4472.66	1997.72	3221.44	2580.77	3953.04
Q12846	Q12846	STX4_HUMAN	STX4				5122.55	4849.13	4309.83	4013.64		3985.14
Q12857	Q12857	NFIA_HUMAN	NFIA	18810.5	25395.3							
Q12860	Q12860	CNTN1_HUMAN	CNTN1					7304.94	4431.07			
Q12874	Q12874	SF3A3_HUMAN	SF3A3				4547.16	5317.37	3844.55	3420.6	4160.66	3942.16
Q12899	Q12899	TRI26_HUMAN	TRIM26		1423.14	2004.57		1605.07	1182.25	1940.95		1953.32
Q12904	Q12904	AIMP1_HUMAN	AIMP1	7133.06	6161.57	5613.3	9384.51	5636.53	6044.36	6530.17	5276.48	6242.34
Q12905	Q12905	ILF2_HUMAN	ILF2	11224	13735.8	9764.98	9129.68	16926.8	12627.6	14310.3	11376	16280.7
Q12906	Q96SI9;Q12906	ILF3_HUMAN	ILF3	10981.6	8294.98	7328.5	10775	14808.6	8568.55	10560.1	6542.47	9833.36
Q12907	Q12907	LMAN2_HUMAN	LMAN2						2227.69	2063.28		2606.01
Q12913	Q12913	PTPRJ_HUMAN	PTPRJ				1908.2	3495.95	4593.49	3367.04	1757.09	3321.79
Q12923	Q12923	PTN13_HUMAN	PTPN13			1366.17	645.045	1995.32	1452.21			
Q12931	Q12931	TRAP1_HUMAN	TRAP1	44957.6	39866.8	43123.2	51880.2	27842.1	39551.1	92282.3	81770.8	106101
Q12959	Q5T0H;Q12959	DLG1_HUMAN	DLG1				3828.52	2211.5	2252.4	3849.95	2460.96	3391.33
Q12965	Q12965	MYO1E_HUMAN	MYO1E	2978.53		3364.64	6686.45	8793.64	7367.8	21207.7	7382.99	17141.7
Q12980	Q12980	NPRL3_HUMAN	NPRL3				2402.86	2291.04	1376.64	3878.93	2267.17	2833.38
Q12982	Q12982	BNIP2_HUMAN	BNIP2					1087.67	2037.37	1025.48		633.709
Q13011	Q13011	ECH1_HUMAN	ECH1	10664.3	7840.67	12425.6	10198.4	3437.05	7743.68	9037.5	11553.1	9300.65
Q13033	Q13033	STRN3_HUMAN	STRN3	5765.42			2354.62	7381.21	5847.05	3238.35		5006.86
Q13043	Q13043	STK4_HUMAN	STK4				3213.16	4666.39	3558.6	4390.73	2924.29	3365.54
Q13045	Q13045	FLII_HUMAN	FLII		3795.21		5444.44	11904.4	10439.7	7890.2	3254.55	5209.78
Q13049	Q13049	TRI32_HUMAN	TRIM32	2838.94	1878.51	2892.54	2346.59	2753.98	2151.33	977.132	1538.37	1067.49
Q13057	Q13057	COASY_HUMAN	COASY				1133.41	2106.8	1825.58	1669.26		3460.46
Q13085	Q13085	ACACA_HUMAN	ACACA	#########	#########	1.73355E+06	2.76387E+06	886726	875101	1.24481E+06	2.91107E+06	1.23865E+06
Q13098	Q13098	CSN1_HUMAN	GPS1	5334.66	3577.75	4775.64	4189.85	1036.18	1825.06	4076.35	2812.75	3129.81
Q13111	Q13111	CAF1A_HUMAN	CHAF1A				1589.14	1207.9	786.172			
Q13112	Q13112	CAF1B_HUMAN	CHAF1B					1420.19		1170.52		
Q13123	Q13123	RED_HUMAN	IK			5965.6	6790.64	13099.5	3680.59	4871.5	5323.38	3896.08
Q13126	Q13126	MTAP_HUMAN	MTAP						4214.76	1019.54		1237.33
Q13131	Q13131	AAPK1_HUMAN	PRKAA1	6385.37	5954.14	7207.45	6755.04	4144.27	6165.74	8024.13	6502.97	8162.35
Q13136	Q13136	LIPA1_HUMAN	PPFIA1	4902.95		4904.12	4624.88	1827.72	1648.34	3853.51	6514.69	4211.03
Q13137	Q13137	CACO2_HUMAN	CALCOCO2		6361.15		3356.4	3448.85	2692.76	7581.89	4155.59	6723.97
Q13144	Q13144	EI2BE_HUMAN	EIF2B5				3520.34	8159.92	4374.38	2293.21	2229.64	6585.37
Q13148	Q13148	TADBP_HUMAN	TARDBP	4998.58	9285.69	5545.3	7073.38	3354.92	4016.89	5677.58	9451.37	6192.87
Q13151	Q13151	ROA0_HUMAN	HNRNPA0	2580.33			2746.32	5309.05	2278.67	2102.1	6039.96	1545.05
Q13153	Q13153	PAK1_HUMAN	PAK1						770.374	1980.38	1633.34	1810.09
Q13155	Q13155	AIMP2_HUMAN	AIMP2	4639.98	5924.26	5787.3	6531.23	9839.65	9457.72	13408.9	8408.78	11645
Q13162	Q13162	PRDX4_HUMAN	PRDX4	4636.44	5385.17	5751.22	10438.5	9484.33	13976.1	21613.1	11975.2	27432.9
Q13177	Q13177	PAK2_HUMAN	PAK2			3650.35	2740.7	2237.61	3954.52	5008.63	5160.4	5111.56
Q13185	Q13185	CBX3_HUMAN	CBX3				891.661	1348.65	2580.17	1171.34	757.766	1302.58
Q13188	Q13188;Q13043	STK3_HUMAN	STK3	4367.21	4199.83	3566.15	4084.26	4139.74	5496.28	2987.75	3101	3156.33
Q13200	Q13200	PSMD2_HUMAN	PSMD2	13549.7	9799.21	17804.3	28478.3	25217.6	32998.8	17696.8	8487.42	17703.9
Q13228	Q13228	SBP1_HUMAN	SELENBP1	7368.58		6891.07	1947.03		1597.44	1380.54		
Q13242	Q13242	SRSF9_HUMAN	SRSF9	18890.2	18974.7	5400.11	3660.1	11544	4724.36	1067.87	5857.26	3702.3
Q13243	Q13243	SRSF5_HUMAN	SRSF5	4972.41		4904.86	4249.43	10199	3584.77	3651.42	8168.16	4210.1
Q13247	Q13247	SRSF6_HUMAN	SRSF6			1506.47	1969.72	932.698	634.902	1095.36	2535.18	1497.07
Q13263	Q13263	TIF1B_HUMAN	TRIM28	5855.84	5114.04	5658.65	11568.7	4272.11	7232.6	11011.2	9368.77	8032.2
Q13277	Q13277	STX3_HUMAN	STX3				4791.17	21981.6	17598.9	15454	5913.72	14926.1
Q13283	Q13283	G3BP1_HUMAN	G3BP1	12855.4			9595.9	37750.5	19222.2	13399	7573.45	15295.3
Q13310	Q13310	PABP4_HUMAN	PABPC4	35432.8	28326.5	9994.38	30621.4	102530	24072.5	14105.7	13116.4	12227
Q13325	Q13325	IFIT5_HUMAN	IFIT5	8912.69	12625.6	10650	3218.55	6159.64	5353.34	3582.12	2116.79	3510.45
Q13330	Q08TCE;Q13330	MTA1_HUMAN	MTA1				8110.22	1780.07	1789.5	4790.56		5571.67
Q13347	Q13347	EIF3I_HUMAN	EIF3I	1931.85		2455.7	3291.52	5046.81	8368	5677.67	2835.52	7680.56
Q13362	Q13362	2A5G_HUMAN	PPP2R5C	705.806		496.496		1898.46	1204.45	2012.94	1609.06	3782.17
Q13363	P56545;Q13363	CTBP1_HUMAN	CTBP1							976.688		1071.47
Q13393	Q13393	PLD1_HUMAN	PLD1		2877.8	2382.73	1327.95	2136.95	2363.96	749.874		958.465
Q13395	Q13395	TARB1_HUMAN	TARBP1				3135.04	2172.86	2393.64	5012.62	4495.89	2659.13
Q13404	Q13404	UB2V1_HUMAN	UBE2V1	15242.4		6267.21	7533.9	3376.91	11566.4	4480.03	4181.94	4277.34
Q13409	Q13409	DC1I2_HUMAN	DYNC1I2	12480.5	12734.1	8456.74	11189.4	6385.59	10074	9861.59	12464.4	8951.49
Q13410	Q13410	BT1A1_HUMAN	BTN1A1				5172.41		1790.96	3560.58	5518.86	3688.35
Q13418	Q13418	ILK_HUMAN	ILK	7457.6	7616.52	9526.97	9870.64	13425.8	12339	12287.9	10885.9	8980.64
Q13421	Q13421	MSLN_HUMAN	MSLN							6125.49	3355.92	4932.04
Q13423	Q13423	NNTM_HUMAN	NNT			1272.34	3366.27		2115.56	2789.11	3934.37	2344.79
Q13425	Q13425	SNTB2_HUMAN	SNTB2	7103.99	5490.84		4275.34	4934.44	8882.21	7321.07	5494.76	5866.81
Q13426	Q13426	XRCC4_HUMAN	XRCC4				1346.06	2316.56	2267.04	1899.92		981.254
Q13428	Q13428	TCOF_HUMAN	TCOF1						1239.33	1131.75	1291.65	1013.1
Q13435	Q13435	SF3B2_HUMAN	SF3B2	12689.9	11010.5	6988.78	9088.33	18784.4	13152.6	11563.2	8440	11432.8
Q13439	Q13439	GOGA4_HUMAN	GOLGA4				23233.3	3723.21	3372.75	2876.97	20885.1	
Q13442	Q13442	HAP28_HUMAN	PDAP1				6831.39	8570.35	3480.43	3898.08	5744.28	5819.61
Q13459	Q13459	MYO9B_HUMAN	MYO9B	1109.35			1010.26	1754.09	1537.86	1386.89	611.116	1553.56
Q13470	Q13470	TNK1_HUMAN	TNK1					1367.45		1635.04	937.306	1818.26
Q13472	Q13472	TOP3A_HUMAN	TOP3A	9155.41	20679.9		6331.19	7311.42	1643.82	2346.68	5678.39	2217.59
Q13488	Q13488	VPP3_HUMAN	TCIRG1				1411.71	3164.78	2058.4	2632.98		1384.46
Q13492	O60641;Q13492	PICAL_HUMAN	PICALM	1442.38	1983.81	3282.73	1895.27	7232.82	7402.29	2513.37	1150.52	3405.01
Q13501	Q13501	SQSTM_HUMAN	SQSTM1	23768.2	23444.3	29563.4	8648.7	7364.69	7702.26	15065.7	8823.13	17720.3
Q13505	Q13505	MTX1_HUMAN	MTX1				6487.86	1185.68	1545.29	2078.57	7915.55	4228.63
Q13509	Q13509	TBB3_HUMAN	TUBB3	25109.7	24695.8	28629.1	9990.55	16497.9	19145.8	31714.8	26423.1	33087.8
Q13526	Q13526	PIN1_HUMAN	PIN1						1119.3			
Q13546	Q13546	RIPK1_HUMAN	RIPK1	3062.6	1153.48	2408.21	3591.01	7356.07	3506.94	2095.97	2817.27	2802.34
Q13547	Q13547;Q92769	HDAC1_HUMAN	HDAC1	5736.25		7488.34	6217.31	6743.06	5060.38	11629.5	6165.49	10112.6
Q13557	Q13557	KCC2D_HUMAN	CAMK2D	8918.68	7189.33	10177.4	9758.18	8881.21	12037.2	11647.8	6708.02	11516.5
Q13561	Q13561	DCTN2_HUMAN	DCTN2	5251	3350.99	4235.06	4174.81	7393.1	11738.8	8224.04	3848.05	6209.68
Q13564	Q13564	ULA1_HUMAN	NAE1				3607.01		3658.54	5146.31	8095.83	4684.2
Q13572	Q13572	ITPK1_HUMAN	ITPK1							3163.52	3094.67	3244.3
Q13573	Q13573	SNW1_HUMAN	SNW1			2052.1	3529.23	4984.07	2985.53	3043.35	5092.77	3273.63
Q13586	Q13586	STIM1_HUMAN	STIM1	2379.55	4200.74	2852.39	1479.66	8852.74	5880.1	4212.14	6148.43	7935.78
Q13595	Q13595	TRA2A_HUMAN	TRA2A				1620.57	2525.94	1575.77			
Q13596	Q13596	SNX1_HUMAN	SNX1	1945.54		1677.13	4735.67	1881.33	1370.71	1325.4	2986.87	1523.06
Q13608	Q13608	PEX6_HUMAN	PEX6	11575.5	6332.56	11804.9	11933.4	4680.69	8974.55	5061.11	1450.98	6059.03
Q13610	Q13610	PWP1_HUMAN	PWP1		2074.29		2453.33	12682.5	7882.93	2018.02	1593.57	1046.56
Q13616	Q13616	CUL1_HUMAN	CUL1	5499.76	7103.66		4292.97	5822.72	4098.31	4128.89	2843.46	4828.14
Q13617	Q13617	CUL2_HUMAN	CUL2	3802.06	1509.54	1944.84	3696.55	3719.24	3496.67	2052.63		1694.74
Q13618	Q13618	CUL3_HUMAN	CUL3			4072.28	1731.75	3050.29	3211.93	2489.45	1626.37	2717.91
Q13619	Q13619	CUL4A_HUMAN	CUL4A	2174.17	2156.13		1966.18	2429.73	1489.03	1326.05	2120.62	1199.85
Q13620	Q13620	CUL4B_HUMAN	CUL4B				918.352	994.332	1022.37	363.668	401.715	754.142
Q13627	Q13627	DYR1A_HUMAN	DYRK1A					5319.96	3110.45	639.842		
Q13636	Q13636	RAB31_HUMAN	RAB31					2849.55		1766.12		2463.04
Q13641	Q13641	TPBG_HUMAN	TPBG	2547.34			2998.44	5187.53	2033.56			
Q13671	Q13671	RIN1_HUMAN	RIN1				11758	3386.21	4146.53	4367.12	7252.05	6881.53
Q13724	Q13724	MOGS_HUMAN	MOGS	8213.33	8052.75	7848.99	8407.91	10920.7	8160.08	8534	4278.63	8396.69
Q13740	Q13740	CD166_HUMAN	ALCAM	2784.14		1559.99	2541.63	8941.71	8197.55	2537.08		1850.67
Q13751	Q13751	LAMB3_HUMAN	LAMB3				3246	3122.02	1954.21	1953.97	1618.51	1928.53
Q13761	Q13761	RUNX3_HUMAN	RUNX3								7931.88	4051.82
Q13813	Q13813	SPTN1_HUMAN	SPTAN1	9371.28	6873.03	10440.7	12946.9	20788.2	16520.1	17388.8	6898.52	14094.3
Q13835	Q13835	PKP1_HUMAN	PKP1	10438.9	6142.15	12631.6	7433.52	5247.57	2078.33	4848.09	4853.22	5456.21
Q13838	Q13838	DX39B_HUMAN	DDX39B	4441.42		1922.18			10115.5	6614.29		
Q13867	Q13867	BLMH_HUMAN	BLMH	51820.5	39775.3	75752.6	35495.3	13357.4	29685.3	36824	25692	42976.2
Q13868	Q13868	EXOS2_HUMAN	EXOSC2				920.637		832.844	984.175	695.499	886.884
Q13884	Q13884	SNTB1_HUMAN	SNTB1						426.583	1258.74	807.724	1055.1
Q13885	Q13885;Q9BVA1	TBB2A_HUMAN	TUBB2A	8870.59	16635.8	8365.5	49183.1	13464.2	14189.9	11979.1	28013.6	17294.1
Q13887	Q13887	KLF5_HUMAN	KLF5	492.944	1149.31	577.816		421.31	295.909	1040.16	312.824	2198.1
Q13895	Q13895	BYST_HUMAN	BYSL				5051.67	6810.68	2839.53	2432.11	2832.34	1673.58
Q13905	Q13905	RPGF1_HUMAN	RAPGEF1	62586.1	55984.4	52172.1		15055.5	11061	39445.2	61465.1	28612.6
Q13950	Q13950	RUNX2_HUMAN	RUNX2	8126.86		10080	8643.80	5074.18	6849.76	9148.87	8790.17	10968.6
Q13951	Q13951	PEBB_HUMAN	CBFB				1543.73		1111.26		2250.93	1729.64
Q14008	Q14008	CKAP5_HUMAN	CKAP5		8780.61	5936.13	9322.51	8966.27	6628.37	6589.55	6326.6	6367.08
Q14011	Q14011	CIRBP_HUMAN	CIRBP				1214.81	5233.69	2891.97	1182.59	1045.95	1083.22
Q14019	Q14019	COTL1_HUMAN	COTL1	3078.8					6672.49	1755.52		1249.12
Q14103	Q14103	HNRPD_HUMAN	HNRNPD	26165.5	18818.5	6228.37	12640.5	19285.5	17526.9	11371.1	4714.82	9033.85
Q14119	Q14119	VEZF1_HUMAN	VEZF1		5447.73			4142.53	1479.91			3665.98
Q14126	Q14126	DSG2_HUMAN	DSG2	19015.3	8314.11	14984.8	18315.9	22605.6	25100	22262.3	10086.4	23423.5
Q14139	Q14139	UBE4A_HUMAN	UBE4A			4670.32	3560.99	3333.24	3458.18	2851.38	3842	3938.44
Q14142	Q14142	TRI14_HUMAN	TRIM14	5693.8		6270.97			1424.85	3530.63		3319.05
Q14152	Q14152	EIF3A_HUMAN	EIF3A	6892.36	9657.65	7427.83	13518.1	13105.1	14672.5	12505.9	8677.64	12544.9
Q14156	Q14156	EFR3A_HUMAN	EFR3A				6471.59	11328.8	9486.17	7656.37	5264.83	8893.47
Q14157	Q14157	UBP2L_HUMAN	UBAP2L	5510.57	3798.03	4309.72	6553.49	7420.81	5265.15	3883.67	3458.86	3589.05
Q14160	Q14160	SCRIB_HUMAN	SCRIB				3516.46	7607.09	5891.04	8948.2	1920.41	5046.37
Q14165	Q14165	MLEC_HUMAN	MLEC			1415.6		1500.09	1225.26			1563.79
Q14166	Q14166	TTL12_HUMAN	TTLL12	7153.52		7609.05	5598.53	5366.89	6180.97	8572.09	4696.55	7006.73
Q14181	Q14181	DPOA2_HUMAN	POLA2					2270.09	4092.02	4840.14	1322.17	4898.39
Q14186	Q14186	TFDP1_HUMAN	TFDP1		657.811		286.879	447.299	299.109	428.85		356.817
Q14191	Q14191	WRN_HUMAN	WRN	29543.8	35863.4		8629.07	12975.5	7384.19	12714.9	8034.64	
Q14192	Q14192	FHL2_HUMAN	FHL2	4502.02	5330.9	6091.56					1800.16	
Q14195	Q14195	DPYL3_HUMAN	DPYSL3	5447.96	6492.52	5479.84	2794.69	1970.77	3150.79	3286.24	1786.85	3019.14
Q14203	Q14203	DCTN1_HUMAN	DCTN1	15576.1	10318.8	10092.5	9748.08	6111.27	8017.05	9082.66	7525.37	10192.6
Q14204	Q14204	DYHC1_HUMAN	DYNC1H1	9728.24	9957.47	7515.51	10046.1	8183.57	7346.09	8351.39	11261.6	8286.14
Q14213	Q14213	IL27B_HUMAN	EBI3	4866.32			1256.99	2198.7	1310.05			
Q14232	Q14232	EI2BA_HUMAN	EIF2B1				4709.18	6123.41	8041.05	5106.31		7034.28
Q14244	Q14244	MAP7_HUMAN	MAP7	12457.5	11827.3	3865.2	10914.2			8567.63	8108.98	12284.1
Q14247	Q14247	SRC8_HUMAN	CTTN	12951.5	12392.7	10043.4	10048.4	9201.21	14934.6	9348.5	8978.78	10258.1
Q14249	Q14249	NUCG_HUMAN	ENDOG		1147.97			1196.83	711.362	2618.11		3364.57
Q14254	Q14254	FLOT2_HUMAN	FLOT2	4709.72	4519.32	4201.8	15985.6	14101.9	14830.5	51076.1	20304.2	40626.3
Q14257	Q14257	RCN2_HUMAN	RCN2	2381.8	5591.74	4904.24	2964.68	5447.4	6214.28	4797.6	3035.91	7906.19
Q14258	Q14258	TRI25_HUMAN	TRIM25	9616.88	8008.3	8875.68	14380.1	18025	16282.8	9394.89	7894.96	8925.89
Q14315	Q14315	FLNC_HUMAN	FLNC	17648.7	18991.1	21177.1	44226	25171.6	30832	5310.76	2445.52	4430.42
Q14318	Q14318	FKBP8_HUMAN	FKBP8	4491.58	4553.67	5354.1	3236.36	1968.57	2167.25	2646.69	3531.93	3468.48
Q14331	Q14331	FRG1_HUMAN	FRG1				5094.21		1833.23	2088.11	5530.42	
Q14344	Q14344	GNA13_HUMAN	GNA13	12581.7	12226.9	16553.1	38548.8	46677.3	58561	214766	48037.3	108019
Q14444	Q14444	CAPR1_HUMAN	CAPRIN1	16202.7	6527.76	8151.05	8891.66	11315.6	5546.26	7798.74	5932.29	8394.89
Q14498	Q14498	RBM39_HUMAN	RBM39	6772.21	4810.14	4561.95	3087.08	4348.08	2676.93	5346.19	4196.94	8083.46
Q14525	Q14525	KT33B_HUMAN	KRT33B	24620.7			4817.4			1731.11	4663.89	4131.32
Q14527	Q14527	HLTF_HUMAN	HLTF	46734.6	67385.8		35080	44577.2	5335.43	15327.2	56061.1	6330.72
Q14532	Q14532	KRT32_HUMAN	KRT32									
Q14558	Q14558	KPRA_HUMAN	PRPSAP1			1729.79	2040.87	2221.77	2068.6	1526.69		1867.25
Q14562	Q14562	DHX8_HUMAN	DHX8			822.519		1269.8		754.905	1928.63	782.138
Q14566	Q14566	MCM6_HUMAN	MCM6			5260.39	2135.7	4898.41	7938.53	8325.64	3449.06	8774.43
Q14571	Q14571	ITPR2_HUMAN	ITPR2					5143.46		1628.3		1325.68
Q14573	Q14573	ITPR3_HUMAN	ITPR3					3252.19	2503.08	6010.71	820.101	5837.49
Q14574	Q14574	DSC3_HUMAN	DSC3	10102.4	7150.25	15583.1	7353.99	1729.9	4079.23	4730.27	6078.49	7078.17
Q14644	Q14644	RASA3_HUMAN	RASA3				449.121	2277.71	2169.13	6732.96	1273.15	4819.41
Q14669	Q14669	TRIPC_HUMAN	TRIP12			1118.07	2675.61	4288.15	4183.36	2313.25	1201.64	2752.63
Q14671	Q14671	PUM1_HUMAN	PUM1	5918.97	2448.71	5215.25	5767.66	6585.73	3796.45	4190.39	6702.37	6290.41
Q14677	Q14677	EPN4_HUMAN	CLINT1	3132.42	2681.61	3125.88	3268.96	3835.78	5462.71	2452.7	1887.17	2404.78
Q14680	Q14680	MELK_HUMAN	MELK				1107.68	3791.62	2149.28	4559.41	2174.8	5912.04
Q14683	Q14683	SMC1A_HUMAN	SMC1A				2782.24	3812	3097.01	4893.08	4668.02	5153.6
Q14684	Q14684	RRP1B_HUMAN	RRP1B	2199.19				2672.5	2753.22	2828.77	1425.69	3246.58
Q14694	Q14694	UBP10_HUMAN	USP10	6126.9	9006.69	8228.78	11708.3	10048.1	6802.98	9068.53	19485.8	10629.6
Q14697	Q14697	GANAB_HUMAN	GANAB	13398.5	10106.1	17737.9	15474.4	17868.1	20831.2	23219.3	9291.75	21362.3
Q14699	Q14699	RFTN1_HUMAN	RFTN1	1622	3090.63	1588.94	5001.02	10580.2	10173.7	2683.9		
Q14730	Q14730	2A5D_HUMAN	PPP2R5D	2118.05	3386.32	3267.13	3858.47	8445.35	6547.9	5986.96	5180.63	5269.2
Q14739	Q14739	LBR_HUMAN	LBR	2207.44	2683.99	1486.98	859.438	2533.68	2490.91	1811.69	2088.06	1559.64
Q14764	Q14764	MVP_HUMAN	MVP	70011	107486	25530.7	26518	27345.9	13614.6	3545.6	4784.56	4281.44
Q14790	Q14790	CASP8_HUMAN	CASP8				1215.5	1132.13	1515.01	709.915		551.975
Q14807	Q14807	KIF22_HUMAN	KIF22				4039.92	3220.95	1580.58	1751.15	2483.62	1268.7
Q14814	Q6Z878;Q14814;Q06413	MEF2D_HUMAN	MEF2D				2435.44	2261.2	1599.85			
Q14839	Q14839;Q8TDI0	CHD4_HUMAN	CHD4				3722.36	6998.61	4432.74	7239.58	5305.39	7504.84
Q14847	Q14847	LASP1_HUMAN	LASP1	9701.14	8597.51	9311.11	9346.74	10380.6	23848.5	7268.24	6925.81	7109.53
Q14914	Q14914	PTGR1_HUMAN	PTGR1		5696.26	6144.2		597.588	2243.44			
Q14919	Q14919	NC2A_HUMAN	DRAP1				3156.87				2743.46	
Q14938	Q14938	NFIX_HUMAN	NFIX	20693.2	20132.1	3934.35						
Q14974	Q14974	IMB1_HUMAN	KPNB1	23318.8	18786.2	21543	36848.8	23021.7	38238.5	29706.5	13013	27742.8
Q14978	Q14978	NOLC1_HUMAN	NOLC1	1415.74	1977.34	1253.96		4046.43	3065.96	5535.96	1606.62	4038.96
Q14997	Q14997	PSME4_HUMAN	PSME4	1592.19			2808.42		768.95		1596.88	672.037
Q14C86	Q14C86	GAPD1_HUMAN	GAPVD1	9642.44	3203.09	8624.93	9656.95	19507.9	8335.27	8027.01	9458.81	6584.79
Q14CN4	Q14CN4	K2C72_HUMAN	KRT72	7229.45		19212.1	6457.28				13485.8	4115.66
Q14CX7	Q14CX7	NAA25_HUMAN	NAA25				1563.1	4576.76	4759.49	823.757		1755.84
Q14CZ7	Q14CZ7	FAKD3_HUMAN	FASTKD3				1401.67		1932.68	6863.37	8779.69	5017.77
Q15003	Q15003	CND2_HUMAN	NCAPH			3812.02	2984.62		2306.49	2621.6	2933.75	2796.1
Q15004	Q15004	PAF15_HUMAN	PCLAF								743.967	
Q15005	Q15005	SPCS2_HUMAN	SPCS2					4951.14	3701.28	7138.4		5363.67
Q15008	Q15008	PSMD6_HUMAN	PSMD6	19811	12503.3	21047.6	25169.1	20685.4	24184.9	20859.3	18449.4	27208.2
Q15014	Q15014	MO4L2_HUMAN	MORF4L2				881.074	2223.79	1623.01	1069.46	1173.03	1241.66
Q15019	Q15019	SEPT2_HUMAN	SEPTIN2	26063.7	18918.5	24717.3	14541.1	15311.5	24743.3	12581.2	5788.3	13799.2
Q15020	Q15020	SART3_HUMAN	SART3	3788.87		1689.71	2898.24	2724.87	2611.47	3326.42	1996.91	2516.32
Q15021	Q15021	CND1_HUMAN	NCAPD2	6290.5	4676.35	3841.06	3944.75	5650.2	5303.28	7758.95	5815.08	8107.23
Q15024	Q15024	EXOS7_HUMAN	EXOSC7					1294.93	1550.72	2253.94		
Q15029	Q15029	U5S1_HUMAN	EFTUD2	4845.32	5626.07	4065.51	6568.61	12165.6	11146.1	14573.2	10380.6	14345.3
Q15042	Q15042	RB3GP_HUMAN	RAB3GAP1	2278.09	3246.19	3681.23	2473.81	1774.24	2269.18	2112.06	1883.3	1917.22
Q15043	Q15043	S39AE_HUMAN	SLC39A14	1337.24			6075.57	2521.53	2489.96	2543.8	3504.87	3292.72
Q15046	Q15046	SYK_HUMAN	KARS1	7058.18		6867.51	5890.13	9625.66	11709.2	13764.4	8431.12	14597.2
Q15054	Q15054	DPOD3_HUMAN	POLD3				1798.57	2921.92	2060.99	2269.25	2719.11	2778.71
Q15056	Q15056	IF4H_HUMAN	EIF4H	9549.81	3888.63	6884.69	3121.15	1156.76	3367.53	2828.38	3263.93	3626.77
Q15067	Q15067	ACOX1_HUMAN	ACOX1				3558.46		3902.54	4665.28	2307.7	4123.58
Q15075	Q15075	EEA1_HUMAN	EEA1	4677.35			2983.09	1708.33	2758.13	1788.72	836.633	2336.19
Q15084	Q15084	PDIA6_HUMAN	PDIA6	11809.2	6412.63	12793.2	10208.3	5440	10319.1	4085.97	4326.6	4788.68
Q15102	Q15102	PA1B3_HUMAN	PAFAH1B3	5306.5		6981.98	6239.77		1734.71	9718.12	3470.73	10057.6
Q15118	Q15118	PDK1_HUMAN	PDK1						1964.18	4373.42	3082.12	
Q15120	Q15120	PDK3_HUMAN	PDK3							10017	5754.52	9517.86
Q15121	Q15121	PEA15_HUMAN	PEA15				2168.49	1471.26	2074.54			
Q15125	Q15125	EBP_HUMAN	EBP					601.745	510.455	1754.75		2391.73
Q15147	Q15147	PLCB4_HUMAN	PLCB4				2554.03	2721.09	2283.23			
Q15149	Q15149	PLEC_HUMAN	PLEC	449939	494086	362861	60442.9	289802	187298	383871	102371	352542
Q15154	Q15154	PCM1_HUMAN	PCM1	6557.07	9031.91	6388.36	4658.79	5267.45	4407.13	3877.22	5128.17	3978.95
Q15165	Q15165	PON2_HUMAN	PON2	4732.32	2151.93	4982.33	4397.71	4301.34	3833.35	5739.69	7388.23	4974.04
Q15172	Q15172	2A5A_HUMAN	PPP2R5A				882.61				665.126	
Q15181	Q15181	IPYR_HUMAN	PPA1				2273.19	2592.55	4131.72	3577.01		3131.21
Q15185	Q15185	TEBP_HUMAN	PTGES3				3313.15	2455.85	5698.73	2755.51	2924.03	4492.59
Q15233	Q15233	NONO_HUMAN	NONO	29230.8	30036.9	28329.1	21684.6	447889	29109.4	59100.8	58232	62022.6
Q15257	Q15257	PTPA_HUMAN	PTPA				1675.23	6338.44	5832.07	2023.47	2116.52	2169.01
Q15274	Q15274	NADC_HUMAN	QPRT	2731.26		2667.49	5322.87	3039.63	11977.1			
Q15286	Q15286	RAB35_HUMAN	RAB35		8766.59	7085.6		28130	39335.4	30383.4	10656.4	30467.4
Q15287	Q15287	RNPS1_HUMAN	RNPS1					9810.73				
Q15293	Q15293	RCN1_HUMAN	RCN1	8097.28	4643.77	8872	17590	8376.56	14440.1	4085.71	8163.86	5823.71
Q15323	O76011;Q14525;O76009;A0A140TA62;Q15323	K1H1_HUMAN	KRT31	13496.4	15878.2	9043.08	10297.8	3757.27	2845.45	5187.09	8655.22	6996.5
Q15334	Q15334	L2GL1_HUMAN	LLGL1				3500.78	3129.86	3792.24	2558.94	3056.66	2540.47
Q15345	Q15345	LRC41_HUMAN	LRRC41				1238.8	3088.69	2791.75	1379.23	410.468	1756.17
Q15365	Q15365	PCBP1_HUMAN	PCBP1	27627.7	16573	33583.6	38972.9	6507.15	15417.3	40167.2	35987.4	33125.5
Q15366	Q15366	PCBP2_HUMAN	PCBP2	59741.5	57668.3	51594.8	91729.9	21021	32342.5	36276.8	76499.3	36494.3
Q15369	Q15369	ELOC_HUMAN	ELOC	5205.17		2786.51	6107.57	3919.69	5140.75	4359.09	3713.35	2383.57
Q15370	Q15370	ELOB_HUMAN	ELOB	4562.97		3956	8958.16	5468.62	11381.3	5568.5	4615.54	6824.59
Q15382	Q15382	RHEB_HUMAN	RHEB	3855.84	5234.64	5809.68	9976.4	4944.6	6509.45	6614.44	5645.52	10221.8
Q15386	Q15386	UBE3C_HUMAN	UBE3C	3208.06		2881.64	3117.54	4214.6	3765.48	5056.13	7011.02	4068.52
Q15392	Q15392	DHC24_HUMAN	DHCR24				2957.45	4737.66	2459.28	3331.47	5614.1	4213.93
Q15393	Q15393	SF3B3_HUMAN	SF3B3	7628.11	7763.37	6625.18	6896.26	8844.65	6229.17	10151	11223.8	10399.6
Q15398	Q15398	DLGP5_HUMAN	DLGAP5				2878.04	4090.29	2121.66	4212.25	2978.24	2296.84
Q15404	Q15404	RSU1_HUMAN	RSU1		4551.47	4665.65	3086.52	11901.1	13756.2	6657.99	2154.71	6421.55
Q15417	Q15417	CNN3_HUMAN	CNN3	6008.46	6882.58	5707.72	9853.59	7182.71	18890.1	2529.63		2317.81
Q15418	Q15349;Q15418;P51812;Q9UK32	KS6A1_HUMAN	RPS6KA1		7443.89	5854.89	5099.79	6533.75	6693.01	9829.56	6350	11926.7
Q15428	Q15428	SF3A2_HUMAN	SF3A2				2073.73	1274.18		2511.06	1679.65	2984.66
Q15434	Q15434	RBMS2_HUMAN	RBMS2	29091.6	10491.3	20233.91	11385.3	8749.68	2954.79	2498.06	3932.78	7959.88
Q15435	Q15435	PP1R7_HUMAN	PPP1R7	6252.62	3487.54	5201.22	3178.34		3475.45	4938.18	1603.98	1881.04
Q15436	Q15436	SC23A_HUMAN	SEC23A	8824.27	9180.51	7104.44	8988	5821.33	5232.3	4806.25	3047.18	4962.98
Q15437	Q15437	SC23B_HUMAN	SEC23B		3101.9		3306.03	4876.89	5138.9	3098.08	2174.79	2937.88
Q15459	Q15459	SF3A1_HUMAN	SF3A1			1527.11	2574.55	5302.19	2872.13	3947.9	2465.88	4753.05
Q15477	Q15477	SKI2_HUMAN	SKIC2	4443.02	4617.87	5687.6	2387.55	5000.31	4248.56	5646.99	2805.88	8438.72
Q15517	Q15517	CDSN_HUMAN	CDSN	26468.7	27037.2	31931.8	17061.2	6901.53	7091.16	15357.1	21450.2	18023.4
Q15554	Q15554	TERF2_HUMAN	TERF2					1675.44	1756	1844.56	762.135	199.987
Q15555	Q15555	MARE2_HUMAN	MAPRE2			3659.5	3495.24	1545.74	2204.33	1735.06	2025.97	3108.35
Q15561	Q15561	TEAD4_HUMAN	TEAD4				1527.23	3201.76	2331.86			
Q15582	Q15582	BGH3_HUMAN	TGFBI	8398.49	8911.47	7408.5		2646.22				5846.78
Q15637	Q15637	SF01_HUMAN	SF1	7906.24	8886.6	6238.71	4616.51	30872.7	9142.82	9287.85	2468.48	8533.57
Q15643	Q15643	TRIPB_HUMAN	TRIP11	2765.26	1982.27		3109.93	1438.98	1548.5	2261.75	2228.66	2207.03
Q15645	Q15645	PCH2_HUMAN	TRIP13	18640.7	19403.5	22123.4	24517.7	21687.8	21132.3	43678.4	35648.6	40488.1
Q15648	Q15648	MED1_HUMAN	MED1		733.228			2423.19	3064.22	2513.72		2957.05
Q15650	Q15650	TRIP4_HUMAN	TRIP4	2837.14			2285.3	4396.58	2113.22	1182.68	2049.98	
Q15654	Q15654	TRIP6_HUMAN	TRIP6	1802.7	2307.77	2079.27	1525.33	4455.59	4062.98			
Q15678	Q15678	PTN14_HUMAN	PTPN14				2122.11	7151.94	4544.05	1735.47		1699.76
Q15691	Q15691	MARE1_HUMAN	MAPRE1	11478.8	3565.8	12741.8	10899.2	9478.93	15551.4	11464.9	4648.75	10037.1
Q15717	Q15717	ELAV1_HUMAN	ELAVL1	4463.99	9066.08	6987.59	3317.75	18188.7	3809.61	3648.93	3258.11	3803.28
Q15723	Q15723	ELF2_HUMAN	ELF2	2729.77	4710.08	3366.55	2073.38	1393.47	872.599		2695.31	1217.28
Q15738	Q15738	NSDHL_HUMAN	NSDHL	4371.78	4120.03	4627.95	3205.81	8257.9	8580.57	15001.4	7918.77	10446.5
Q15758	Q15758	AAAT_HUMAN	SLC1A5	8458.92	10755.1	11185.6	12958.5	33749.1	30528	79730.1	43822.5	48578.8
Q15773	Q15773	MLF2_HUMAN	MLF2			2094.5		2921.37	3168.72	4311.49	4573.91	6401.72
Q15785	Q15785	TOM34_HUMAN	TOMM34			1259.84	3091.13	4215.65	8721.33	4629.28	1407.42	3015.48
Q15796	O15198;Q15796;P84022	SMAD2_HUMAN	SMAD2	5974.05	9905.91	5808.45	3926.28	1348.11	2991.16	4190.17	2611.46	3452.39
Q15813	Q15813	TBCE_HUMAN	TBCE				2999.53	2062.35	4894.91	2226.99	1761.79	2657.86
Q15833	Q15833	STXB2_HUMAN	STXBP2				664.684	1439.36	1180.03	2524.81	954.189	2260.84
Q15836	P23763;Q15836;P63027	VAMP3_HUMAN	VAMP3	2388.12					1452.03	905.419		1602.18
Q15904	Q15904	VAS1_HUMAN	ATP6AP1				4178.05		1897.65	2508.71	3211.21	1531.16
Q15907	P62491;Q15907	RB11B_HUMAN	RAB11B	6101.11	4056.91	8081.32	9336.93			10337.5	11362.3	9531.4
Q15942	Q15942	ZYX_HUMAN	ZYX	10592.3	6030.48	9315.21	6082.75	2565.35	5513.98	1998.36	3655.65	2351.34
Q16181	Q16181;Q6ZU15	SEPT7_HUMAN	SEPTIN7	9564.97	7200.96	9929.8	13137.9	6427.5	12465.6	6683.63	4723.16	5285.75
Q16186	Q16186	ADRM1_HUMAN	ADRM1				2158.1	1827.55	3965.6	4642.81	9737.46	3990.07
Q16204	Q16204	CCDC6_HUMAN	CCDC6					2982.01	3570.91	675.764		
Q16222	Q16222	UAP1_HUMAN	UAP1	3350.83		4259.75	2023.11	1903.1	5929.63	1420.29	790.788	1255.65
Q16254	Q16254	E2F4_HUMAN	E2F4							616.477	1404.66	1072.84
Q16352	Q16352	AINX_HUMAN	INA	96622.8	132018	97774.1	3824.75	4954.62	21459.2	13603.7	5952.13	29293.8
Q16378	Q16378	PROL4_HUMAN	PRR4								1831.01	
Q16401	Q16401	PSMD5_HUMAN	PSMD5	5500.44	4911.81	4795.5	8678.02	6358	6540.49	4137.34	4000.61	3049.91
Q16512	Q16512	PKN1_HUMAN	PKN1				1817.34	8594.71	6810.6	4774.82	2458.92	1451.44
Q16513	Q16513	PKN2_HUMAN	PKN2			1449.24	2136.81	1720.13	1945.03	1505.9	1552.93	1202.16
Q16515	Q16515	ASIC2_HUMAN	ASIC2	6075.65	4320.78	20617	2625.57		1762.61	1007.08	3537.95	8120.31
Q16531	Q16531	DDB1_HUMAN	DDB1	7273.49	9343.08	5779.76	7677.44	16231.4	12468.7	7118.35	8036	8634.01
Q16543	Q16543	CDC37_HUMAN	CDC37	3220.8			4395	2524.33	5276.91	2914.13	1947.75	4637.68
Q16555	Q16555	DPYL2_HUMAN	DPYSL2	8796.35	5661.12	8566.25	3997.54	3049.98	3558.18	4019.37	4429.27	4158.52
Q16563	Q16563	SYPL1_HUMAN	SYPL1	4879.22	4764.86	5726.08	2752.79	16883.5	20615.4	10298.4	1723.67	13677.4
Q16576	Q16576	RBBP7_HUMAN	RBBP7	8855.32	9281.39	5904.02	10623.6	13565.2	9641.64	12070.8	10330	14445.6

Q16595	Q16595	FRDA_HUMAN	FXN									
Q16610	Q16610	ECM1_HUMAN	ECM1	6902.1	4695.18	7458.99	3644.9	1458.9	1753.51	2112.92	3910.18	3052.64
Q16629	Q16629	SRSF7_HUMAN	SRSF7	1510.57	1538.67		784.084	4432.54	1060.24	747.34	1373.64	1041.32
Q16630	Q16630	CPSF6_HUMAN	CPSF6	5949.35	11356	2687.45	3980.22	66825.1	7272.97	5189.27	3281.51	4969.88
Q16637	Q16637	SMN_HUMAN	SMN2				2913.8	10589	4913.59	4334.04	7368.36	5646.21
Q16643	Q16643	DREB_HUMAN	DBN1	4486.53		3538.96	3556.28	11407	7464.91	25044.6	4251.76	10706.1
Q16656	Q16656	NRF1_HUMAN	NRF1	173674	81273		3231.27	22727.2	42361.3		15437.5	
Q16658	Q16658	FSCN1_HUMAN	FSCN1	5869.92		8758.54	5548.81	4376.68	10873.4	5155.36		4618.26
Q16666	Q16666	IF16_HUMAN	IFI16	7626.38	5478.4	5586.62	4173.25	8654.32	5789.89			
Q16678	Q16678	CP1B1_HUMAN	CYP1B1									
Q16698	Q16698	DECR_HUMAN	DECR1	8517.32	11074	10394	4184.46	4036.98	2373.09	2834.95	5797.47	3624.92
Q16718	Q16718	NDUA5_HUMAN	NDUFA5				560.427	741.025	570.467	1832.11	844.296	1090.3
Q16719	Q16719	KYNU_HUMAN	KYNU	2657.3	794.026	3982.24						
Q16739	Q16739	CEGT_HUMAN	UGCG				1420.88	1737.05	2440.97			
Q16762	Q16762	THTR_HUMAN	TST							2226.51		2633.47
Q16795	Q16795	NDUA9_HUMAN	NDUFA9					5276.65	3631.01	15547.6	6656.72	13381.7
Q16822	Q16822	PCKGM_HUMAN	PCK2	7683.67	6555.19	7488.63	5478.02	1530.48	2933.34	3254.65	4691.46	4192.56
Q16850	Q16850	CP51A_HUMAN	CYP51A1			3140.03		2263.19	4445.62	2965.79	1775.44	4034.28
Q16851	Q16851	UGPA_HUMAN	UGP2	5892.55		5918.7	7312.43	4268.79	3892.35	2970.71	3859.27	2905.14
Q16881	Q9NNW7;Q16881	TRXR1_HUMAN	TXNRD1	5399.99	4229.17	7420.18	2988.63	4298.19	9356.45	2142.4	1165	2244.49
Q16891	Q16891	MIC60_HUMAN	IMMT	34587	44493.8	28817.1	3709.97	11675.4	7369.75	7304.1	2763.15	9084.53
Q1KMD3	Q1KMD3	HNRL2_HUMAN	HNRNPUL2	3188.42		4035.03	1532	3275.22	2746.87	3696.41	2385.32	3442.45
Q27J81	Q27J81	INF2_HUMAN	INF2	12351.1	8803.23	8228.38	4260.43	6724.16	3155.45	6300.73	7968.63	9108.3
Q29RF7	Q29RF7	PDS5A_HUMAN	PDS5A	6008.8	6071.83	4073.04	6216.94	5420.25	5010.27	10712.8	11682.8	7721.34
Q2KHR3	Q2KHR3	QSER1_HUMAN	QSER1				1312.6	1033.79	635.219	1626.08	1546.06	1331
Q2M1P5	Q2M1P5	KIF7_HUMAN	KIF7					14389	10088.9			
Q2M3D2	Q2M3D2	EX3L2_HUMAN	EXOC3L2				5401.07	8734.59	5490.96			
Q2NKX8	Q2NKX8	ERC6L_HUMAN	ERCC6L				1735.77	3744.03	2517.98	4181.07	1913.22	3842.98
Q2NL82	Q2NL82	TSR1_HUMAN	TSR1				10953.3	28229.4	13893.1	11094.3	9579.59	9749.36
Q2TAL8	Q2TAL8	QRIC1_HUMAN	QRICH1			1843.63	1530.29	1282.99	1751.55	2371.87		1670.55
Q2TAY7	Q2TAY7	SMU1_HUMAN	SMU1			2609.84	3222.74		3058.45	6853.68		1949.31
Q2TB90	Q2TB90	HKDC1_HUMAN	HKDC1							9514.43	17978.6	7837.1
Q2V2M9	Q2V2M9	FHOD3_HUMAN	FHOD3	27466	22980.2		6173.83	7417.21				
Q2VPK5	Q2VPK5	CTU2_HUMAN	CTU2				3486.68			4639.2	7096.27	3379.81
Q32MZ4	Q32MZ4	LRRF1_HUMAN	LRRFIP1				2882.38		2200.77	1156	2823.29	1451.4
Q32NB8	Q32NB8	PGPS1_HUMAN	PGS1	2009.38	824.863	1569.4	1721.89	2459.77	2091.59	2613.33	1513.98	3405.62
Q32P28	Q32P28	P3H1_HUMAN	P3H1				1851.38	3339.9	3114.16	1067.61		2379.39
Q3KQU3	Q3KQU3	MA7D1_HUMAN	MAP7D1	4914.63	5721.24	3167.03	3800.89	6579.47	5658.78	4091.76	3155.26	2867.55
Q3LXA3	Q3LXA3	TKFC_HUMAN	TKFC			2197.49	1507.74	1348.6	2693.14	3807.62	2648.49	2973.12
Q3MHD2	Q3MHD2	LSM12_HUMAN	LSM 12:00				1996.22	6839.32	2393.32	1048.79		1160.56
Q3MIP1	Q3MIP1	IPIL2_HUMAN	ITPRIPL2								303.163	
Q3MIT2	Q3MIT2	PUS10_HUMAN	PUS10									
Q3SXY8	Q3SXY8	AR13B_HUMAN	ARL13B							2801.78	970.003	2877.63
Q3SY69	Q3SY69	AL1L2_HUMAN	ALDH1L2	8325.08	6410.32	7300.15	9637.82	3019.58	2749.4		1178.08	
Q3SY84	Q3SY84	K2C71_HUMAN	KRT71	2633.28		3953.76	4281.26	1530.31	3040.97		1371.14	5018.12
Q3V6T2	A6NC98;Q3V6T2	GRDN_HUMAN	CCDC88A	4283.38			2638.52	2351.11	2134.73	3821.99	2200.17	4268.93
Q3ZCQ8	Q3ZCQ8	TIM50_HUMAN	TIMM50	17122.5	26598.3	16148.4	24350.8	9964.65	9326.66	14128.5	31758.9	15777.8
Q460N5	Q460N5	PAR14_HUMAN	PARP14	6725.53	5262.54	3659.51	2449.19	3298.02	2024.02	1841.19	6834.15	1504.66
Q4G0T1	Q4G0T1	SRCRM_HUMAN	SCART1					14606.5	23061.8	23977.4		38107.5
Q4KMP7	Q4KMP7	TB10B_HUMAN	TBC1D10B		1513.23			4450.28	2161.71	2926.76		
Q4KMQ2	Q4KMQ2	ANO6_HUMAN	ANO6			726.745		1372.76	1129.82	1252.47		1038.33
Q4V339;Q5JTY5;Q5R1A9;Q8IUF	Q8IUF1;Q5R1A9;Q9BRT8;Q4V339;Q5JTY5	ZNG1A_HUMAN;ZNG1B_HUMAN;ZNG1C_H	ZNG1A;ZNG1B;ZNG1C;ZNG1E;ZNG1F						2224.18	2203.41	2199.37	2096.29
Q52LJ0	Q52LJ0	FA98B_HUMAN	FAM98B	3583.34		2315.64	2595.13	6167.52	5775.54	5592.31	2721.17	4589.67
Q53EP0	Q53EP0	FND3B_HUMAN	FNDC3B	8434.27	7727.33	6523.55	4337.94	3268.08	4627.32	1815.5		2453.64
Q53EU6	Q53EU6	GPAT3_HUMAN	GPAT3				3110.44	2815.5	2884.21	1317.92	1760.57	2260.3
Q53EZ4	Q53EZ4	CEP55_HUMAN	CEP55					5159.96	3430.81	6647.7	2289.39	6226.3
Q53GQ0	Q53GQ0	DHB12_HUMAN	HSD17B12	11543.6	14739.7	10352.4	11522.9	14166	11698.5	22726.1	22182.7	23293.4
Q53H12	Q53H12	AGK_HUMAN	AGK	4441.49	3735.6	2944.15				2840.9	1768.94	3638.57
Q53H96	Q53H96	P5CR3_HUMAN	PYCR3								1077.84	
Q53RT3	Q53RT3	APRV1_HUMAN	ASPRV1	4484.63	4102.09	4840.64	2742.72			1037.17	7019.63	
Q53S08;Q9H0N0	Q9H0N0;Q53S08	RAB6C_HUMAN;RAB6D_HUMAN	RAB6C;RAB6D					913.836	1071.52	1674.61	1165.67	1007.92
Q562E7	Q562E7	WDR81_HUMAN	WDR81	2010.85			1668.33	1550.04	1569.99	3417.39	2356.55	2972.78
Q562R1	Q562R1	ACTBL_HUMAN	ACTBL2	253340	345610	219603	352152	616459	196834	405363	315619	308957
Q56VL3	Q56VL3	OCAD2_HUMAN	OCIAD2							2289.47	830.376	3092.38
Q58EX7	Q58EX7	PKHG4_HUMAN	PLEKHG4			1288.13	1082.74	2160.34	1158.62	1561.72	1636.08	
Q58FF3	Q58FF3	ENPLL_HUMAN	HSP90B2P									658.577
Q58FF8	Q58FF8	H90B2_HUMAN	HSP90AB2P					2855.61	4957.03	1284.91	902.889	924.576
Q5BJD5	Q5BJD5	TM41B_HUMAN	TMEM41B				1392.5	1380.05	1056.33	4621.43	2199.58	5129.06
Q5D862	Q5D862	FILA2_HUMAN	FLG2	29485.8	24246.7	45204.9	23064.4	5436.24	13688.2	23880.3	19312	29121.9
Q5EB52	Q5EB52	MEST_HUMAN	MEST								5505.72	
Q5EBL8	Q5EBL8	PDZ11_HUMAN	PDZD11	2604.91	5479.45	1976.17	843.28	4005.04	4451.3	1735.14		1894.95
Q5GLZ8	Q5GLZ8	HERC4_HUMAN	HERC4	5858.56	8178.12	5994.21	1184.07	1173.1	1391.36	2568.23	778.686	2134.09
Q5H9R7	Q5H9R7	PP6R3_HUMAN	PPP6R3		2168.02		3884.29	4625.5	7324.27	3678.9	3430.84	5053.3
Q5HYK3	Q5HYK3	COQ5_HUMAN	COQ5					3231.47	2842.85	3532.59	2448.32	4599.91
Q5JPH6	Q5JPH6	SYEM_HUMAN	EARS2	3546.02	5050.58	4676.64	2275.8	2881.52	2916.83	3405.68	2680.12	4179.27
Q5JRX3	Q5JRX3	PREP_HUMAN	PITRM1	675.549		1182.06	768.044		866.966			
Q5JSZ5	Q5JSZ5	PRC2B_HUMAN	PRRC2B	3361.12			1714.32	5369.43	3409.28	843.877		3299.63
Q5JTH9	Q5JTH9	RRP12_HUMAN	RRP12				2056.63	3747.29	3330.73	1045.29	1490.1	2421.84
Q5JTV8	Q5JTV8	TOIP1_HUMAN	TOR1AIP1	2643.14				2566.69	3822.62	5230.35	2215.53	4537.08
Q5JTZ9	Q5JTZ9	SYAM_HUMAN	AARS2		2558.74	3891.87	3324.51	2123.11	2762.63	4564.26	4379.18	3410.97
Q5JVF3	Q5JVF3	PCID2_HUMAN	PCID2		2178.24		644.374	3570.06	2074.7	835.427	977.809	867.235
Q5JWF2	Q5JWF2	GNAS1_HUMAN	GNAS	6213.72		11421	14118.8	24273.9	23096.2	47326.1	11251.4	42144.4
Q5M775	Q5M775	CYTSB_HUMAN	SPECC1				2411.29	5466.76	1938.51	2039.38	342.518	2153.33
Q5SRE5	Q5SRE5	NU188_HUMAN	NUP188				3964.11	2800.03	3186.5	3040.9	5472.92	2982.21
Q5SSJ5	Q5SSJ5	HP1B3_HUMAN	HP1BP3	6498.99	5739.07	3924.18		2024.48	2059.95		2160.84	
Q5ST30	Q5ST30	SYVM_HUMAN	VARS2						829.921	1297.06		941.997
Q5SW79	Q5SW79	CE170_HUMAN	CEP170	2900.06	3048.06	2064.33	3897.93	4822.03	2865.85	1201.93	1456.89	841.941
Q5SY16	Q5SY16	NOL9_HUMAN	NOL9					3088.6	949.291	870.276		
Q5SZB4	Q5SZB4	CI050_HUMAN	C9orf50							6352.27		
Q5T0B9	Q5T0B9	ZN362_HUMAN	ZNF362						11797.7	3043.3		
Q5T160	Q5T160	SYRM_HUMAN	RARS2	4496.71	4615.82	3959.71	2300.47	3237.59	2476.41	2894.53	2578.22	3847.49
Q5T1M5	Q5T1M5	FKB15_HUMAN	FKBP15			371.748	1605.33	1689.82	1109.42	1058.21	1294.96	917.579
Q5T447	Q5T447	HECD3_HUMAN	HECTD3				1946.17	3333.63	1270.68	2004.73	1218.98	2748.18
Q5T4S7	Q5T4S7	UBR4_HUMAN	UBR4		3451.54	5366.83	4645.71	2728.22	2576.9	3868.97	5612.47	3463.96
Q5T5P2	Q5T5P2	SKT_HUMAN	KIAA1217	6287.44			5876.46	7782.34	5816.8	3530.38	2261.45	4173.12
Q5T6F2	Q5T6F2	UBAP2_HUMAN	UBAP2				4614.12	2281.65	2747.28	2990.86	3557.85	2862.62
Q5T6V5	Q5T6V5	QNG1_HUMAN	QNG1				1862.11		1532.17			
Q5T749	Q5T749	KPRP_HUMAN	KPRP	77636.2	61107.7	115034	58446.5	21719.1	30793	52918.8	33718.2	70201.1
Q5T750	Q5T750	XP32_HUMAN	XP32	22945.4	17123.3	35608.7	16588.6	4842	4207.7	9083.27	13187.7	18911
Q5T8P6	Q5T8P6	RBM26_HUMAN	RBM26	1659.75	1480.2	1279.37	1453.28	2887.98	1134.02	1489.09	2253.14	1545.19
Q5T9A4	Q5T9A4	ATD3B_HUMAN	ATAD3B	6868.18	8679.38	4064.13		2406.2		7310.45	1286.88	7959.64
Q5TA45	Q5TA45	INT11_HUMAN	INTS11				1541.86	5377.83	3672.69	5629.11	6397.38	5420.48
Q5TEC6	Q5TEC6;P84243;P68431;Q71DI3;Q16695;Q6NXT2	H37_HUMAN	H3-7	312529	311259	193871	154288	291448	150696	53868	83331.4	42303.4
Q5UIP0	Q5UIP0	RIF1_HUMAN	RIF1					627.9	588.634	3058.57	1157.42	1332.25
Q5VIR6	Q5VIR6	VPS53_HUMAN	VPS53	2434.67		1598.34	2336.86	2134.94	796.637	2234.33	3713.75	2251.09
Q5VSL9	Q5VSL9	STRP1_HUMAN	STRIP1				2340.59	2348.4	2428.19	2275.52	1586.21	3781.3
Q5VT25	Q5VT25	MRCKA_HUMAN	CDC42BPA				2240.2	3066.41	2257.31	2704.61	2417.72	1901.2
Q5VUA4	Q5VUA4	ZN318_HUMAN	ZNF318		824.345			1201.97	674.224	985.352		774.128
Q5VV42	Q5VV42	CDKAL_HUMAN	CDKAL1							1591.59	1882.76	1522.92
Q5VYK3	Q5VYK3	ECM29_HUMAN	ECPAS	6905.9	6231.87	6273.87	10871.6	7460.79	7908.87	9107.54	9464.73	9125.64
Q5VYS8	Q5VYS8	TUT7_HUMAN	TUT7	3558.34		1784.42	1470.05			1319.25		2147.56
Q5VZE5	Q5VZE5	NAA35_HUMAN	NAA35				1276.13	1914.81	1135.9	1249.63	712.605	1353.2
Q5XPI4	Q5XPI4	RN123_HUMAN	RNF123						1129.92	802.336		1017.79
Q63HN8	Q63HN8	RN213_HUMAN	RNF213			1947.1	2023.9	1504.18	1493.37	1271.16	2351.67	1753.4
Q63ZY3	Q63ZY3	KANK2_HUMAN	KANK2	9898.61	10086.3	8195.13	5509.12	4508.63	3953.45	2964.83	4699.01	3671.4
Q643R3	Q643R3	LPCT4_HUMAN	LPCAT4		1199.74		1320.04	9295.88	3690.62	2877.07	1090.79	8322.57
Q658Y4	Q658Y4	F91A1_HUMAN	FAM91A1				2190.37		1505.37	2147.15	2328.24	2736.31
Q66K74	Q66K74	MAP1S_HUMAN	MAP1S	2830.02	1989.22	2563.87	3306.94	1972.48	3392.31	2840.83	1765.65	3221.04
Q68CQ7	Q68CQ7	GL8D1_HUMAN	GLT8D1		2747.42	1541.55		1994.29	1268.34	2625.29	787.491	2598.76
Q68CZ2	Q63HR2;Q68CZ2;Q9HBL0	TENS3_HUMAN	TNS3			5923	3408.17	6107.66	1896.12	9293.78	9526.27	9005.7
Q69YL0	Q69YL0	NCAS2_HUMAN	NCBP2AS2					3482.39	665.376	837.621		
Q69YQ0	Q69YQ0	CYTSA_HUMAN	SPECC1L	6087.56		8660.76	4567.99	26529.8	7924.16	2973.79		3458.71
Q6AI08	Q6AI08	HEAT6_HUMAN	HEATR6					2504.09	1848.46	696.358		
Q6AWC2	Q6AWC2	WWC2_HUMAN	WWC2				718.264	1251.06	996.543			750.278
Q6DD88	Q6DD88	ATLA3_HUMAN	ATL3	9668.57	9273.45	9956.62	8997	7915.64	6252.4	2964.4	3451.41	2475.48
Q6DKJ4	Q6DKJ4	NXN_HUMAN	NXN				732.985	1774.48	1817.83	739.081		2062.2
Q6IA86	Q6IA86	ELP2_HUMAN	ELP2					1739.61	2059.54	1231.83		740.842
Q6IAA8	Q6IAA8	LTOR1_HUMAN	LAMTOR1	4666.38	5343.49	3883.43	10041.8	16810.1	21773.9	49458.8	17307	39105.7
Q6IBS0	Q6IBS0	TWF2_HUMAN	TWF2				1310.14	2471.75	1985.34	1137.02		1470.58
Q6IQ23	Q6IQ23	PKHA7_HUMAN	PLEKHA7				1554.74	8024.45	4336.75	5604.94	4782.34	7411.63
Q6JQN1	Q6JQN1	ACD10_HUMAN	ACAD10					996.127	1185.15	2112.12	1496.09	2356.06
Q6KB66	Q6KB66	K2C80_HUMAN	KRT80	9185.45	12801.9	17743.7	11462.4	41415.7	24811.2	41256.3	15663.1	41248.9
Q6L8Q7	Q6L8Q7	PDE12_HUMAN	PDE12				2444.51		1647.82	1119.65	1313.96	1843.8
Q6MZP7	Q6MZP7	LIN54_HUMAN	LIN54	1382.95			647.987	3594.96	1012.57			
Q6NSI4	Q6NSI4	RADX_HUMAN	RADX								2283.79	
Q6NSJ5	Q6NSJ5	LRC8E_HUMAN	LRRC8E				1389.3	2701.92	2038.53	1006.66		1090.21
Q6NTF9	Q6NTF9	RHBD2_HUMAN	RHBDD2							1332.33	418.442	1463.17
Q6NUM9	Q6NUM9	RETST_HUMAN	RETSAT	1882.32		1590.31	1993.75	1773.04	1728.53	1270.81	1525.09	1518.42
Q6NUQ4	Q6NUQ4	TM214_HUMAN	TMEM214	5771.96	5435.72	3262.61	3572.47			2096.43	3243.77	2169.95
Q6NXE6	Q6NXE6	ARMC6_HUMAN	ARMC6	4578.77	4129.83	4280.37	7363.33	5794.46	5638.77	7344.42	9830.37	8272.39
Q6NXG1	Q6NXG1	ESRP1_HUMAN	ESRP1							2900.45	4442.03	
Q6NXR4	Q6NXR4	TTI2_HUMAN	TTI2				2914.99	1363.29	857.093	1389.11	2230.55	2281.86
Q6NYC8	Q6NYC8	PPR18_HUMAN	PPP1R18	3973.21	4765.14	2509.79		4739.45	3883.12	3979.11		
Q6NZ67;Q6P582	Q6NZ67;Q6P582	MZT2A_HUMAN;MZT2B_HUMAN	MZT2A;MZT2B					847.783	720.746			419.804
Q6NZI2	Q6NZI2	CAVN1_HUMAN	CAVIN1	8572.08	6088.42	8004.55	12096.4	12702.8	11773.1	11338.6	6217.5	10054.3
Q6NZY4	Q6NZY4	ZCHC8_HUMAN	ZCCHC8					1385.36	834.949	679.489		
Q6P087	Q6P087	RUSD3_HUMAN	RPUSD3		897.619			1016.96	1042.3	1988.34	1085.45	1639.98
Q6P158	Q6P158	DHX57_HUMAN	DHX57					6513.28	1888.53			
Q6P1J9	Q6P1J9	CDC73_HUMAN	CDC73					2178.5	1578.3	1506.08	980.828	1071.97
Q6P1L8	Q6P1L8	RM14_HUMAN	MRPL14	4537.11	3216.21	5279.27	3572.14	2111.81	1963.2	2928.58	6412.33	2557.99
Q6P1M3	Q6P1M3	L2GL2_HUMAN	LLGL2							1999.79	1026.93	1225.93
Q6P1N0	Q6P1N0	C2D1A_HUMAN	CC2D1A	2691.26	3371.8	2527.41	2747.57	1562.86	1779.06	1928.4	951.812	1891
Q6P2C8	Q6P2C8	MED27_HUMAN	MED27				1433.4	1871.52	1743.21	1079.01		411.554
Q6P2E9	Q6P2E9	EDC4_HUMAN	EDC4	22886.6			4412.29	11550.9	6862.37	9280.43	6307.01	7837.55
Q6P2H3	Q6P2H3	CEP85_HUMAN	CEP85		2068.16		1943.78	1818.67	1963.24	4050.92	1414.9	3835.95
Q6P2Q9	Q6P2Q9	PRP8_HUMAN	PRPF8		5820.41		5397.65	12024.9	6834.61	8336.47	10045.8	7453.88
Q6P3W7	Q6P3W7	SCYL2_HUMAN	SCYL2			1566.76	1499.75	2726.31	2893.94	1412.42	1155.08	1564.08
Q6P3X3	Q6P3X3	TTC27_HUMAN	TTC27				3602.53	7436.85	4930.63	6714.43	2828.52	5289.67
Q6P4A8	Q6P4A8	PLBL1_HUMAN	PLBD1	3983.83		5799.3	2701.49		2766.08	1874.97	3410.82	2554.86
Q6P5R6	Q6P5R6	RL22L_HUMAN	RPL22L1	7387.61	9907.79	9486.25	22332.2	53413.8	31519.6	2001.83	7404.05	1160.63
Q6P996	Q6P996	PDXD1_HUMAN	PDXDC1				6192.8	6361.22	3771.14	4814.99	9131.06	5338.57
Q6P9B6	Q6P9B6	MEAK7_HUMAN	MEAK7				922.188	6965.47	7177.71	2485.71		2281.19
Q6P9B9	Q6P9B9	INT5_HUMAN	INTS5				4952.84	2196.99	1153.5	2614.29	8179.95	
Q6PD62	Q6PD62	CTR9_HUMAN	CTR9				1094.36	2041.98	1688.65	1953.67	985.389	1581.94
Q6PD74	Q6PD74	AAGAB_HUMAN	AAGAB					2876.04	3938.84	3690.71		733.35
Q6PGP7	Q6PGP7	SKI3_HUMAN	SKIC3	4370.6	4543.4	4380.83	4622.22	6068.5	4063.44	4673.18	4531.87	3115.5
Q6PI48	Q6PI48	SYDM_HUMAN	DARS2	2063.91	2967.25	1406.95	1401.64	4752	7309.35	2844.13	2591	2722.62
Q6PIU2	Q6PIU2	NCEH1_HUMAN	NCEH1	6614.92	3790.44	6229.78	4088.33	2784.33	2848.18	7782.68	6590.1	7703.48
Q6PJG2	Q6PJG2	MDEAS_HUMAN	MIDEAS					4809.44	2130.72	1266.09		
Q6PJG6	Q6PJG6	BRAT1_HUMAN	BRAT1				3275.54	2354.73	1159.45	2840.37	2729.55	3935.06
Q6PKG0	Q6PKG0;Q659C4	LARP1_HUMAN	LARP1	12856.9	14866.2	7958.06	8826.25	48714.4	15134.2	10337.3	6661.7	9596.08
Q6PUV4	Q6PUV4	CPLX2_HUMAN	CPLX2									
Q6QEF8	Q6QEF8	CORO6_HUMAN	CORO6				2864.93	7016.39	4112.31			
Q6R327	Q6R327	RICTR_HUMAN	RICTOR	2130.78	2187.22	1119.57	2194.54	2123.13	2020.05	2528.83	2230.86	3266.83
Q6RFH5	Q6RFH5	WDR74_HUMAN	WDR74					2781.76	1494.65	3210.13	2012.72	2890.93
Q6UB35	Q6UB35	C1TM_HUMAN	MTHFD1L			7471.2	8119.3		8918.56	12219.5	12576.1	12735.7
Q6UN15	Q6UN15	FIP1_HUMAN	FIP1L1	7692.02		5458.04	5144.11	3653.08	3064.92	3320.27	7852.71	3316.79
Q6UWJ1	Q6UWJ1	TMCO3_HUMAN	TMCO3			8344.07		13505.6		13585	8179.15	13767.8
Q6UWP8	Q6UWP8	SBSN_HUMAN	SBSN	8877.94	6517.15	17106.4	7028.25	10852.2	7190.95	6950.24	8536.71	9344.76
Q6UX53	Q6UX53	TMT1B_HUMAN	TMT1B						843.656	3807.25	4978.53	4164.05
Q6VY07	Q6VY07	PACS1_HUMAN	PACS1				759.599	1825.49	2417.2	256.211	1097.89	612.338
Q6W2J9	Q6W2J9	BCOR_HUMAN	BCOR									
Q6WCQ1	Q6WCQ1	MPRIP_HUMAN	MPRIP	24606	24919.5	20798.9	73066.7	91229.8	61013.6	10568.3	10225.7	9930.99
Q6XE24	Q6XE24	RBMS3_HUMAN	RBMS3	43829.9	30291.1		12126.8	8820.68	740.817			
Q6XQN6	Q6XQN6	PNCB_HUMAN	NAPRT	1551.09					993.947	1484.8	2265.03	965.49
Q6Y7W6	Q6Y7W6	GGYF2_HUMAN	GIGYF2	3964.96	3438.81	3900.01	3891.72	5080.71	3395.83	2584.11	3567.47	2616.85
Q6YHK3	Q6YHK3	CD109_HUMAN	CD109	3584.49		1798.74	2368.17	2574.9	2127.4	2560.74	2064.37	3588.85
Q6YN16	Q6YN16	HSDL2_HUMAN	HSDL2			1759.17	4346.42	10951.7	5226.68	4240.45	5127.67	4139.57
Q6ZMP0	Q6ZMP0	THSD4_HUMAN	THSD4									
Q6ZN17	Q6ZN17	LN28B_HUMAN	LIN28B				9821.75	23841.9	17240.9	3721.42		
Q6ZN28	Q6ZN28	MACC1_HUMAN	MACC1					1157.94		2451.84	1211.24	2119.57
Q6ZNB6	Q6ZNB6	NFXL1_HUMAN	NFXL1	826.397	974.039		1134.64	352.106	511.682	2151.43	1684.26	2152.28
Q6ZRV2	Q6ZRV2	FA83H_HUMAN	FAM83H	2129.1	6487.83	2782.75	1541.13	6299.46	5815.7	13475.1	2661.06	14676.9
Q6ZS17	Q6ZS17	RIPR1_HUMAN	RIPOR1	2583.35			4249.87	5128.94	5674.87	4472.57	2466.99	3902.53
Q6ZSR9	Q6ZSR9	YJ005_HUMAN						1544.63	1281.35			
Q6ZTW0	Q6ZTW0	TPGS1_HUMAN	TPGS1							2618.86		
Q6ZVD8	Q6ZVD8	PHLP2_HUMAN	PHLPP2	353768	342339	268583	303835	82266.4	84578.1	160183	311727	144702
Q6ZVX7	Q6ZVX7	FBX50_HUMAN	NCCRP1	10028.6		13367.8	7613.91		4557.96	5522.72	9749.15	9002.19
Q6ZW31	Q6ZW31	SYDE1_HUMAN	SYDE1	3153.81	2297.01	2071.21	2726.33					
Q70CQ2	Q70CQ2	UBP34_HUMAN	USP34	79996.8	77070.4	99183.8	47174.4	7751.29	8199.19	13219.9	64596.5	19535.4
Q70J99	Q70J99	UN13D_HUMAN	UNC13D				9717.14	10670.7	11497			689.093
Q70SY1	Q70SY1	CR3L2_HUMAN	CREB3L2									
Q71RC2	Q71RC2	LARP4_HUMAN	LARP4	3404.22	4808.78		3659.74	12041.8	6441.11	4413.86	5345.08	6004.3
Q71SY5	Q71SY5	MED25_HUMAN	MED25	2892.9		2893.83	3805.12	793.723	1423.78	3125.79	4591.23	3452.62
Q71UM5	Q71UM5	RS27L_HUMAN	RPS27L	10308.7	11906.5	10310.4	16307.5	16542.3	12284.5	7234.31	10770.4	7944.41
Q75N03	Q75N03	HAKAI_HUMAN	CBLL1	1484.26	2320.18	2095.42		3345.91	2342.86	2930.07	804.511	2782.31
Q75QN2	Q75QN2	INT8_HUMAN	INTS8								1902.7	
Q7KZF4	Q7KZF4	SND1_HUMAN	SND1	7848.39	4625.85	8195.99	12853.4	13979.3	16696.6	12955.3	6988.35	11220.2
Q7KZI7	Q7KZI7	MARK2_HUMAN	MARK2				1659.02	2563	1480.79	3144.79	2676.1	2593.66
Q7L014	Q7L014	DDX46_HUMAN	DDX46	4892.41	4730.05	2423.61	3701.73	3295.51	2881.04	2934.73		3465.48
Q7L0Y3	Q7L0Y3	TM10C_HUMAN	TRMT10C				2482.09	3906.47	3132.59	2681.4	1624.66	3912.57
Q7L1Q6	Q7L1Q6	BZW1_HUMAN	BZW1	11341.6	996.127	7047.39	2755.82	3059.79	5722.53	3297.93	3939.92	4046.4
Q7L2E3	Q7L2E3	DHX30_HUMAN	DHX30	6788.23	6845.87	6235.93	4526.76	12856.1	6730.82	6397.96	4951.97	4781.32
Q7L2H7	Q7L2H7	EIF3M_HUMAN	EIF3M				1985.52	6092.36	7759.36	5175.54	1284.69	5796.02
Q7L3T8	Q7L3T8	SYPM_HUMAN	PARS2		1577.95		1262.95	3378.81	1898.18	3136.4	2509.23	2448.89
Q7L576	Q7L576	CYFP1_HUMAN	CYFIP1	5986.96	8639.23	4084.74	8240.49	8494.92	8073.74	4478.39	4554.8	3519.49
Q7L592	Q7L592	NDUF7_HUMAN	NDUFAF7	3061.48			2042.14			777.242	5696.75	990.325
Q7L5N1	Q7L5N1	CSN6_HUMAN	COPS6	957.011		1122.55	2317.97		2881.4	2918.28		1945.66
Q7L8L6	Q7L8L6	FAKD5_HUMAN	FASTKD5				6833.16	5072.97	3243.26	11001.6	10638	10477.4
Q7L9B9	Q7L9B9	EEPD1_HUMAN	EEPD1				2114.2	10855.2	3751.52	45667.1	14256.7	48524.2
Q7RTP6	Q7RTP6	MICA3_HUMAN	MICAL3				1413.91	14186.1	6163.31	1767.32		
Q7RTS7	Q7RTS7	K2C74_HUMAN	KRT74	25951.3	18892.6	44866.9	20314.9	77799.3	57327.3	40582.8	31202	43299.3
Q7Z2E3	Q7Z2E3	APTX_HUMAN	APTX		2744.84		2019.53	4237.91	1072			844.975
Q7Z2K8	Q7Z2K8	GRIN1_HUMAN	GPRIN1				2266.79	2808.52	2620.64	8059.28	3478.55	7055.84
Q7Z2W4	Q7Z2W4	ZCCHV_HUMAN	ZC3HAV1	4647.92	6495.95	7543.55	13355.2	10314.7	5932.32	3475.03	6086.85	4089.13
Q7Z2Y5	Q7Z2Y5	NRK_HUMAN	NRK				1410.91	3133.69	2177.87			
Q7Z3B4	Q7Z3B4	NUP54_HUMAN	NUP54				2846.21			781.607	455.952	
Q7Z3U7	Q7Z3U7	MON2_HUMAN	MON2	5888.56		4171.01	7772.8	7049.36	6110.87	5864.5	4014.72	5389.33
Q7Z3Y8	Q7Z3Y8;Q7Z3Z0	K1C27_HUMAN	KRT27			8514.06	4036.33		4780.49		2376.76	6810.2
Q7Z417	Q7Z417	NUFP2_HUMAN	NUFIP2	6670.45		6586.11	11526.5	16536.1	4952.31	3360.94	5444.38	4476.61
Q7Z422	Q7Z422	SZRD1_HUMAN	SZRD1						672.779	1104.26	713.362	1214.97
Q7Z434	Q7Z434	MAVS_HUMAN	MAVS	3158.21	3155.55	3047.35		928.066	2207.28	2040.2		2986.05
Q7Z460	Q7Z460	CLAP1_HUMAN	CLASP1			899.726	998.526	876.793	1100.23	1124.56	2726.61	1089.53
Q7Z478	Q7Z478	DHX29_HUMAN	DHX29			1078.54		3260.94	1685.51	1118.97		
Q7Z494	Q7Z494	NPHP3_HUMAN	NPHP3					1964.01	1229.28			
Q7Z4H7	Q7Z4H7	HAUS6_HUMAN	HAUS6				4624.85	2481.99	2159.89	2131.93	1802.47	2200.87
Q7Z4H8	Q7Z4H8	PLGT3_HUMAN	POGLUT3					3782.76	3286.9			
Q7Z4Q2	Q7Z4Q2	HEAT3_HUMAN	HEATR3	3375.75	4184.87	2916.98	8363.45	13766.5	6079.71	11512.1	14221.2	8947.13
Q7Z4S6	Q7Z4S6	KI21A_HUMAN	KIF21A				686.089	1171.06	1017.03	1443.76	826.982	1152.38
Q7Z4W1	Q7Z4W1	DCXR_HUMAN	DCXR			8183.51	2307.85	838.313	7060.85	8651.54	6982.59	8783.91
Q7Z5G4	Q7Z5G4	GOGA7_HUMAN	GOLGA7			1983.88			3814.34	12684.9	2735.41	7746.39
Q7Z5L9	Q7Z5L9	I2BP2_HUMAN	IRF2BP2							2182.95	1624.93	2168.35
Q7Z628	Q7Z628	ARHG8_HUMAN	NET1							1222.13		2041.44
Q7Z6I8	Q7Z6I8	CE024_HUMAN	C5orf24				3204	5514.08	2659			
Q7Z6J8	Q7Z6J8	UBE3D_HUMAN	UBE3D						584.115	1164.78		
Q7Z6W7	Q7Z6W7	DNJB7_HUMAN	DNAJB7	377625	234649	240279	10579	45632.5	27096.5	36076.3	14736.5	44844.6
Q7Z6Z7	Q7Z6Z7	HUWE1_HUMAN	HUWE1				3360.98	2044.5	987.293	4759.07	4637.26	5213.38
Q7Z739	Q7Z739;Q9BYJ9	YTHD3_HUMAN	YTHDF3				3465.11	8626.15	4275.96	2652.38	3203.77	3740.52
Q7Z794	Q7Z794	K2C1B_HUMAN	KRT77	25925.4	25257	47215.6	20537.7	5209.98	12044	6420.3	21645.2	27009.6
Q7Z7A3	Q7Z7A3	CTU1_HUMAN	CTU1				11895	9406.41	3429.89	5474.45	9925.69	7883.32
Q86SE5	Q86SE5	RALYL_HUMAN	RALYL				6937.98	1879.93	977.066	2446.89	6840.13	5279.71
Q86SQ0	Q86SQ0	PHLB2_HUMAN	PHLDB2	6876.1	8172.13	4888.39	4180.36	8338.83	4168.06		1344.52	
Q86T24	Q86T24	KAISO_HUMAN	ZBTB33		847.944					608.099	1270.59	1024.15
Q86TG7	Q86TG7	PEG10_HUMAN	PEG10				863.847	2599.65	1027.57	2107.86	3567.24	1890.09
Q86TV6	Q86TV6	TTC7B_HUMAN	TTC7B				1849.28	3378.99	3214.07	5083.85		5231.89
Q86U38	Q86U38	NOP9_HUMAN	NOP9		6323.69		1687.63	5721.61		2444.61	1408.49	2767.14
Q86U90	Q86U90	YRDC_HUMAN	YRDC				3322.21	2174.17	2203.54	1892.02		1911.42
Q86UE4	Q86UE4	LYRIC_HUMAN	MTDH	5125.41	4561.82	4411.85	3463.51	2893.84	3399.76	5675.06	3039.06	5798.92
Q86UE8;Q9UKI8	Q9UKI8;Q86UE8	TLK1_HUMAN;TLK2_HUMAN	TLK1;TLK2					1075.39	791.529	477.968		510.292
Q86UK7	Q86UK7	ZN598_HUMAN	ZNF598					1227	1300.97			
Q86UP2	Q86UP2	KTN1_HUMAN	KTN1	3942.46			2913.76	3695.44	2976.5	3358.59	1804.43	3214.66
Q86US8	Q86US8	EST1A_HUMAN	SMG6					1103.12		1261.43		
Q86UU1	Q86UU1	PHLB3_HUMAN	PHLDB3		721.049		1163.07	2701.62	1479.59	894.071	532.11	987.094
Q86UW9	Q86UW9	DTX2_HUMAN	DTX2					1151.37	598.35	974.857		1018.86
Q86UY6	Q86UY6	NAA40_HUMAN	NAA40					2633.27	2385.59	2344.69		1837.6
Q86V48	Q86V48	LUZP1_HUMAN	LUZP1		3923.41		3163.79	5397.87	4449.19			1455.68
Q86V81	Q86V81	THOC4_HUMAN	ALYREF	13247.8	10723.6	16305.6	18721.7	37303.2	25709.4	21673.5	16047.8	17244.2
Q86VI3	Q86VI3	IQGA3_HUMAN	IQGAP3	1610.37	4898.2		6352.28	6155.46	4763.01	3931.7	3754.96	4586.86
Q86VN1	Q86VN1	VPS36_HUMAN	VPS36	3014.16		2312.58	2681.64	1633.12	1175.94	1330.64	497.521	1317.73
Q86VP6	Q86VP6	CAND1_HUMAN	CAND1	18063.1	10345.5	17084.2	35326.8	30383.2	27164.1	26984.7	19877.8	25164.2
Q86VR2	Q86VR2	RETR3_HUMAN	RETREG3				1204.73	4146.48	2843.59	2362.71		
Q86VS8	Q86VS8	HOOK3_HUMAN	HOOK3	55451.4	34071.5	37887.2	49904.3	18590.9	14483.9	16057.1	32775.6	15634.9
Q86VY4	Q86VY4	TSYL5_HUMAN	TSPYL5									
Q86W42	Q86W42	THOC6_HUMAN	THOC6					1332.69	637.236	412.977		891.903
Q86W92	Q86W92;Q8ND30	LIPB1_HUMAN	PPFIBP1			5500.38	6633.25	5429.7	4074.14	5642.98	5208.53	6511.83
Q86WB0	Q86WB0	ZC3C1_HUMAN	ZC3HC1				410.296	595.148	452.047	938.278	375.723	832.069
Q86WJ1	Q86WJ1	CHD1L_HUMAN	CHD1L	7866.87	13199.1	3695.88	7113.58	9516.72	4104.41	5061.06	6009.8	4787.78
Q86X29	Q86X29	LSR_HUMAN	LSR				1904.44	1616.18	3329.31	20380.7	4882.89	14748
Q86X55	Q86X55	CARM1_HUMAN	CARM1	3342.97		2818.47	1592.7	1817.05	1558.52	3053.8	3171.35	2617.42
Q86XI2	Q86XI2	CNDG2_HUMAN	NCAPG2		1299.64		1902.08	2057.84	1207.98	4099.33	3029	4054.7
Q86XP3	Q86XP3	DDX42_HUMAN	DDX42				1907.64	6841.94	7674.99	7357.24	696.788	7049.66
Q86XZ4	Q86XZ4	SPAS2_HUMAN	SPATS2				1505.71	1247.07	1208.16	1822.69	1652.97	1992.64
Q86Y07	Q86Y07	VRK2_HUMAN	VRK2		2595.08			2110.97	1222.47	2061.72		897.006
Q86Y39	Q86Y39	NDUAB_HUMAN	NDUFA11	2958.02	2447.07		1925.69			3683.18	1304.11	1411.2
Q86Y56	Q86Y56	DAAF5_HUMAN	DNAAF5	8525.37	4107.82		8405.4	6816.73	7482.92	14359.6	9627.91	15155.8
Q86Y79	Q86Y79	PTH_HUMAN	PTRH1	5464.7	5693	5303.12				1615.76		3017.49
Q86Y82	Q86Y82	STX12_HUMAN	STX12	3974.6	2792.39	2903.61	1408.11	2175.15	3698.55	2969.33		2388.01
Q86YP4	Q86YP4	P66A_HUMAN	GATAD2A				2422.12	521.557	636.415	1131.96	2114.93	890.054
Q86YQ8	Q9HCH3;Q86YQ8;Q8IYJ1	CPNE8_HUMAN	CPNE8				458.648	2089.46	1986.29			
Q86YT6	Q86YT6	MIB1_HUMAN	MIB1					2365.93	1763.08	1559.48		1907.08
Q86YV9	Q86YV9	HPS6_HUMAN	HPS6						1978.29	6117.03	5004.04	4239.9
Q86YW0	Q86YW0	PLCZ1_HUMAN	PLCZ1		7834.25	3788.86		6811.06	2875.55		1977.93	3849.44
Q86YZ3	Q86YZ3	HORN_HUMAN	HRNR	151845	95918.7	187715	136718	37387	46333.6	71166.1	83917.2	113881
Q8IUC1	Q8IUC1	KR111_HUMAN	KRTAP11-1									
Q8IUC4	Q8IUC4;A8MT19	RHPN2_HUMAN	RHPN2		10337.8		7109.89	6768.26	5828.89			
Q8IUD2	Q8IUD2	RB6I2_HUMAN	ERC1	3340.56	3677.21	2157.16	2928.39	3167.28	2209.7	1046.39		879.386
Q8IUE6	Q96QV6;P16104;Q8IUE6	H2A2B_HUMAN	H2AC21	10263.1	12870.6	9085.55	11487.1	8072.09	5816.92			
Q8IUR7	Q8IUR7	ARMC8_HUMAN	ARMC8				5771.17			1121.13	5317.59	2118.91
Q8IUX4	Q8IUX4	ABC3F_HUMAN	APOBEC3F		2715.39			4335.2	1694.7			2369.88
Q8IV08	Q8IV08	PLD3_HUMAN	PLD3		5230.1			1993.34	2208.67		733.045	237.641
Q8IV36	Q8IV36	HID1_HUMAN	HID1									
Q8IV63	Q8IV63	VRK3_HUMAN	VRK3		4231.29		10187.2	4087.24	3782.67	1079.4	2870.06	
Q8IVF2	Q8IVF2	AHNK2_HUMAN	AHNAK2					1226				
Q8IVF7	Q8IVF7	FMNL3_HUMAN	FMNL3	3348.84		1987.15	6351.96	4825.62	4933.17	3993.49	6432.31	3131.28
Q8IVL6	Q8IVL6	P3H3_HUMAN	P3H3		1576.71	1267.32	729.434	5744.24	4755.5	1432.59	1024.27	
Q8IVM0	Q8IVM0	CCD50_HUMAN	CCDC50						2164.35			
Q8IVS2	Q8IVS2	FABD_HUMAN	MCAT			2817.68	1138.11		994.41	3474.58	2765.03	3018.11
Q8IVT2	Q8IVT2	MISP_HUMAN	MISP		11313.7	5834.63			1800.13	12147.3	3091.14	9146.63
Q8IVT5	Q8IVT5	KSR1_HUMAN	KSR1					2311.27				
Q8IVW6	Q8IVW6	ARI3B_HUMAN	ARID3B				698.674	7082.07	2030.85			
Q8IW19	Q8IW19	APLF_HUMAN	APLF	8757.43	10318.8		15229.3					
Q8IW75	Q8IW75	SPA12_HUMAN	SERPINA12	4719.72	3843.61	5380.7	3195.71	2482.82	2919.35	2325.35		3060.53
Q8IWE2	Q8IWE2	NXP20_HUMAN	FAM114A1	2420.44	1189.32	1795.03	536.992	926.722	1797.18			
Q8IWE4	Q8IWE4	DCNL3_HUMAN	DCUN1D3					3987.95	3245.74	7567.52	7457.75	8959.72
Q8IWN7	Q8IWN7	RP1L1_HUMAN	RP1L1	7351.67	5497.41	8435.39	13715.4		2375.01	8685.96	17786.9	9452.28
Q8IWS0	Q8IWS0	PHF6_HUMAN	PHF6					7068.83	5273.46	12141.9	7042.03	14247
Q8IWT6	Q8IWT6	LRC8A_HUMAN	LRRC8A				1158.11	1939.56	3427.93	3963.73	1799.84	2166.01
Q8IWU2	Q8IWU2	LMTK2_HUMAN	LMTK2				1938.89	3773.27	1916.15	5617.76	1774.79	5037.8
Q8IWV7	Q8IWV7	UBR1_HUMAN	UBR1	853.718	3709.44		1464.99	1780.39	2698.44	689.74		790.943
Q8IWX8	Q8IWX8	CHERP_HUMAN	CHERP				2333.51	4741.36	2421.28	2528.57	2593.72	2261.64
Q8IWZ3	Q8IWZ3	ANKH1_HUMAN	ANKHD1				6246.41	5180.11	2495.73	1488.38	5923.18	1902.37
Q8IX12	Q8IX12	CCAR1_HUMAN	CCAR1				1298.57	849.946	533.991	828.732	795.632	674.259
Q8IXB1	Q8IXB1	DJC10_HUMAN	DNAJC10				2222.74	2273.86	1855.8	2781.52	1497.27	2704.69
Q8IXH7	Q8IXH7	NELFD_HUMAN	NELFCD					4966.6	2534.6	5090.46		9076.33
Q8IXM3	Q8IXM3	RM41_HUMAN	MRPL41		5316.56		2566		2636.05	3584.86	2355.02	
Q8IXS6	Q8IXS6;C9JA33	PALM2_HUMAN	PALM2					1891.79	1710.74	16334	2675.75	14901.8
Q8IXS8	Q8IXS8	HYCC2_HUMAN	HYCC2							3518.73		3379.32
Q8IY17	Q8IY17	PLPL6_HUMAN	PNPLA6	4122.5	5548.87	3902.72	2391.64	6496.27	3903.13	5952.96	2821.05	6882.93
Q8IY21	Q8IY21	DDX60_HUMAN	DDX60				867.557	2035.16	2482.98			
Q8IY22	Q8IY22	CMIP_HUMAN	CMIP				1780.28		1927.59	6061.59	3368.49	6918.93
Q8IY67	Q8IY67	RAVR1_HUMAN	RAVER1	4305.72	8563.77	5647.33	7585.96	6121.95	10939.4	11142.3	4738.43	12561.3
Q8IY81	Q8IY81	SPB1_HUMAN	FTSJ3					5540.81	1430.24	482.178		447.487
Q8IYB3	Q8IYB3	SRRM1_HUMAN	SRRM1	1693.36	2869.4	569.35	813.957	4658.04	1292.96	1393.88	1567.6	925.813
Q8IYB8	Q8IYB8	SUV3_HUMAN	SUPV3L1				1978.16	1267.19	2038.16	2474.06	2327.4	2874.15
Q8IYI6	Q8IYI6	EXOC8_HUMAN	EXOC8		951.188	1591.27	1460.14	1370.84	1225.32	1238.81	965.54	1233.11
Q8IYT3	Q8IYT3	CC170_HUMAN	CCDC170				35041.4	29635	68161.5	263411	24380.9	196565
Q8IZ52	Q8IZ52	CHSS2_HUMAN	CHPF	3649.17	1285.72	2715.36	2126.33	1623.8	1234.4	9462.96	6643.1	11968

Q 8IZ69	Q8IZ69	TRM2A_HUMAN	TRMT2A				5987.43	4621.76	1513.46	758.987	7742.79	1695.87
Q 8IZ83	Q8IZ83	A16A1_HUMAN	ALDH16A1	1352.1	1583.96	919.365	2676.58	1719.64	1785.95	1092.91	440.733	1403.23
Q 8IZH2	Q8IZH2	XRN1_HUMAN	XRN1	13767.7	21168.1		5327.68	5157.91	3822.9	3210.31	2355.28	4374.59
Q 8IZL8	Q8IZL8	PELP1_HUMAN	PELP1			1220.94	5263.82	9749.89	3387.96	1275.05	495.65	1490.37
Q 8IZP0	Q9NYB9;Q8IZP0	ABI1_HUMAN	ABI1			5774.49	3504.28	9759.97	7713.26	6438.74		6724.27
Q 8IZV5	Q8IZV5	RDH10_HUMAN	RDH10	1840.1	1887.98	1717.62			672.599	4885.73	2827.63	5093.49
Q 8IZW8	Q8IZW8	TENS4_HUMAN	TNS4							3769.86	1577	3594.89
Q 8N0X7	Q8N0X7	SPART_HUMAN	SPART				1851.64	2169.77	1611.63			
Q 8N0Z3	Q8N0Z3	SPICE_HUMAN	SPICE1					1627				
Q 8N122	Q8N122	RPTOR_HUMAN	RPTOR		3575.42	2811.79	4283.87	2490.14	2383.79	4796.94	6540.5	4563.73
Q 8N163	Q8N163	CCAR2_HUMAN	CCAR2	2934.52	2796.94	4968.22	5211.39	11134.2	10811.3	16901.7	9361.12	12799.9
Q 8N183	Q8N183	NDUF2_HUMAN	NDUFAF2				2608.55			7315.53	5191.24	4551.22
Q 8N1B4	Q8N1B4	VPS52_HUMAN	VPS52			2206.37	2315.11		1294.1	2083.62	764.753	2405.23
Q 8N1F7	Q8N1F7	NUP93_HUMAN	NUP93		5938.96	4663.87	12206	10112.7	7755.15	14787.8	13982.3	14967.7
Q 8N1G4	Q8N1G4	LRC47_HUMAN	LRRC47	4526.4	3024.59	4076.82	5113.66	5413.28	7751.86	3327.62	2564.06	2784
Q 8N1N4	Q8N1N4	K2C78_HUMAN	KRT78	87681.1	65749.8	114765	69888.1	31504.8	35897.4	44951.5	68818	63458.5
Q 8N201	Q8N201	INT1_HUMAN	INTS1	3608.29		2732.67	4119.17	11744.5	2622.63	4786.53	7689.8	4668.83
Q 8N283	Q8N283	ANR35_HUMAN	ANKRD35				2868.35			3302.93		
Q 8N335	Q8N335	GPD1L_HUMAN	GPD1L					1925.59				
Q 8N392	Q8N392	RHG18_HUMAN	ARHGAP18				986.804	1512.55	2058.6	4718.73		1866.7
Q 8N3C0	Q8N3C0	ASCC3_HUMAN	ASCC3	15249.9	11532.4	2167.36	13266.6	11357.2	6257	6327.21	13750.4	6136.76
Q 8N3E9	Q8N3E9	PLCD3_HUMAN	PLCD3				3902.8	3113.27	3241.74	11852.7	5158.7	12681.2
Q 8N3R9	Q8N3R9	PALS1_HUMAN	PALS1					3385.82	1330.13	1770.4		1470.89
Q 8N3V7	Q8N3V7	SYNPO_HUMAN	SYNPO	14359.7	6748.5	12266.4						
Q 8N488	Q8N488	RYBP_HUMAN	RYBP				1884.63					
Q 8N4C8	Q8N4C8	MINK1_HUMAN	MINK1			1791.69	897.074	1151.75	1622.38	1117.38	2045.93	1744.88
Q 8N4V1	Q8N4V1	EMC5_HUMAN	MMGT1	3486.91	3260.87	2472.51	969.818	1204.08	1509.04	5540.81	1645.12	5822.23
Q 8N4X5	Q8N4X5	AF1L2_HUMAN	AFAP1L2				2153.48	2604.32	2958.8			
Q 8N556	Q8N556	AFAP1_HUMAN	AFAP1	1540.47		2215.23	1330.83	591.248	709.148	1039.6	1024.86	755.256
Q 8N5K1	Q8N5K1	CISD2_HUMAN	CISD2		5679.76			4867.53	4965.45	7764.92		9323.86
Q 8N5M9	Q8N5M9	JAGN1_HUMAN	JAGN1					2946.92	3109.92	3672.05		3516.67
Q 8N680	Q8N680	ZBTB2_HUMAN	ZBTB2		4327.71			567.074				
Q 8N684	Q8N684	CPSF7_HUMAN	CPSF7				9182.58	69175.8	12024.1	6472.91	7158.36	5642.93
Q 8N6L1	Q8N6L1	KTAP2_HUMAN	KRTCAP2	2014.01			966		427.57	1468.08	1858.05	772.029
Q 8N6R0	Q8N6R0	EFNMT_HUMAN	METTL13		6643.68	5462.05	6925.41	5725.96	5381.29	5800.64	7357.96	5220.17
Q 8N6T3	Q8N6T3	ARFG1_HUMAN	ARFGAP1				5203.44	3085.14	4081.09	1244.39		1199.38
Q 8N766	Q8N766	EMC1_HUMAN	EMC1				1052.62	1195.84	1170.49	1679.97	1204.18	1529
Q 8N7H5	Q8N7H5	PAF1_HUMAN	PAF1				2339.7	3526.07	4457.55	2675.6	1949.28	
Q 8N7R7	Q8N7R7	CCYL1_HUMAN	CCNYL1				4760.87	5909.67	4104.95	6721.1	6269.42	5386.11
Q 8N884	Q8N884	CGAS_HUMAN	CGAS		935.565		7315.28	30899.8	16396.8			
Q 8N8S7	Q8N8S7	ENAH_HUMAN	ENAH	1433.42		1448.27	378.378	1154.35	1894.21	287.632		201.297
Q 8N9B5	Q8N9B5	JMY_HUMAN	JMY					1737.85	1572.04	2704.07		1188.44
Q 8N9N2	Q8N9N2	ASCC1_HUMAN	ASCC1	8802.17	4012.18		3829.46	2363.1	1117.28	1239.12	3843.6	
Q 8NB37	Q8NB37	GALD1_HUMAN	GATD1						1483.84	793.192		
Q 8NB90	Q8NB90	SPAT5_HUMAN	SPATA5		3694.32		4567.31	4027.35	2754.49	2142.62	2429.64	2168.6
Q 8NBF2	Q8NBF2	NHLC2_HUMAN	NHLRC2			2739.18	1715.98	2890.39	3266.15	1651.07		1440.96
Q 8NBJ5	Q8NBJ5	GT251_HUMAN	COLGALT1				3663.86	5070.75	6109.18			
Q 8NBJ7	Q8NBJ7	SUMF2_HUMAN	SUMF2	3845.18	1591.12	2805.29	2950.82	1734.52	2259.34	1769.45	1057.38	2213.89
Q 8NBN7	Q8NBN7	RDH13_HUMAN	RDH13				830.59	1211.85	1423.13	1686.33	780.258	1403.45
Q 8NBQ5	Q8NBQ5	DHB11_HUMAN	HSD17B11	1386.39	2473.12	1697.84	604.674	3520.74	3986.25	5113.19	1504.15	1769.7
Q 8NBS9	Q8NBS9	TXND5_HUMAN	TXNDC5	8018.17		6709.22	2934.1	14058	21182.4	7669.89	2831.27	6089.66
Q 8NBU5	Q8NBU5	ATAD1_HUMAN	ATAD1					4964.87	3256.84	2485.26		2637.47
Q 8NBX0	Q8NBX0	SCPDL_HUMAN	SCCPDH	4382.64	3049.43	5987.33	2716.22	6511.84	6534.59	5367.76	2120.64	4698.95
Q 8WUK0	Q8WUK0	PTPM1_HUMAN	PTPMT1	182061	183231	209192		3535.01	2371.11	6836.34	4148.06	8139.1
Q 8WUM0	Q8WUM0	NU133_HUMAN	NUP133				1566.27	3275.16	2504.46	1966.88		1030.24
Q 8WUM4	Q8WUM4	PDC6I_HUMAN	PDCD6IP	11874.1	9236.45	14337.6	10527.1	8548.57	11079.7	11834.2	6984.84	9048.22
Q 8WUW1	Q8WUW1	BRK1_HUMAN	BRK1				843.725	675.656	1531.45		456.159	451.022
Q 8WV24	Q8WV24	PHLA1_HUMAN	PHLDA1			3061.7	1727.66		1448.81	4578.53	2907.21	4041.78
Q 8WV41	Q8WV41	SNX33_HUMAN	SNX33	2463.31	4029.18	1947.75	1891.13		1897.49			1138.72
Q 8WV74	Q8WV74	NUDT8_HUMAN	NUDT8								1149.46	1396.3
Q 8WVB6	Q8WVB6	CTF18_HUMAN	CHTF18		317.922		1617.98	3298.68	2141.5	3248.41	1791.72	1997.15
Q 8WVC6	Q8WVC6	DCAKD_HUMAN	DCAKD			5834.62	3339.67	1959.15	2348.6	3157.43	4309.92	4672.45
Q 8WVM8	Q8WVM8	SCFD1_HUMAN	SCFD1	5022.46	5083.19	3518.42	4922.33	2693.98	3369.74	2774.83	2305.56	3392.76
Q 8WVV4	Q8WVV4	POF1B_HUMAN	POF1B	5215.99		4232.61	3691.81	2252.24	2052.03	14310.7	8153.25	12556.4
Q 8WVV9	Q8WVV9	HNRLL_HUMAN	HNRNPLL					3359.46	3080.96	3077.57		2839.51
Q 8WVX9	Q8WVX9	FACR1_HUMAN	FAR1	3368.77			3362.51			4813.17	6869.37	3085.19
Q 8WVY7	Q8WVY7	UBCP1_HUMAN	UBLCP1					1163.5	1679.09	375.019		
Q 8WW12	Q8WW12	PCNP_HUMAN	PCNP	1452.57			930.059	799.357	2083.89	938.901	588.645	843.773
Q 8WW33	Q8WW33	GTSF1_HUMAN	GTSF1				5198.31		6576.24			
Q 8WWI1	Q8WWI1	LMO7_HUMAN	LMO7	13580.8	15186.7	14485.1	2309.28	6883.65	2733.93	6986.48	11481	7518.63
Q 8WWI5	Q8WWI5	CTL1_HUMAN	SLC44A1					2986.49	3963.43	1195.42		790.468
Q 8WWM7	Q8WWM7	ATX2L_HUMAN	ATXN2L	7156.31	6890	5738.01	13298.3	46726.8	14401.8	7011.2	5374.76	7277.76
Q 8WWZ7	Q8WWZ7	ABCA5_HUMAN	ABCA5				1733.17	3739.52		4060.59		4159.4
Q 8WX92	Q8WX92	NELFB_HUMAN	NELFB				1516.87	1857.34	2418.52	860.317	566.261	1831.29
Q 8WX93	Q8WX93	PALLD_HUMAN	PALLD	4075.88	4151.13	2864.73	2874.3	4827.76	5382.2			
Q 8WX94;Q 9NX02	Q9NX02;Q8WX94	NALP2_HUMAN;NALP7_HUMAN	NLRP2;NLRP7									
Q 8WXE1	Q8WXE1	ATRIP_HUMAN	ATRIP				1864.58				4733.77	
Q 8WXE9	Q8WXE9	STON2_HUMAN	STON2	1566.7		2683.28	162725	18672.9	52109.2	552312	120842	476457
Q 8WXF1	Q8WXF1	PSPC1_HUMAN	PSPC1	2785.43		3795.7	2285.31	66261.1	9213.68	4369.15	2170.42	4647.86
Q 8WXG6	Q8WXG6	MADD_HUMAN	MADD				421.71		1407.97	1421.18		1777.47
Q 8WXI7	Q8WXI7	MUC16_HUMAN	MUC16		2400.8			11897.8	14931.5	8848.2	2800.62	3167.16
Q 8WXI9	Q8WXI9	P66B_HUMAN	GATAD2B	743.805	680.06	229.814	2641.81		345.282	1228.11	2209.49	1451.06
Q 8WXR4	Q8WXR4	MYO3B_HUMAN	MYO3B		23425.4	9110.06	16412.6	6785.49	9685.75	55268.5	8179.4	
Q 8WXW3	Q8WXW3	PIBF1_HUMAN	PIBF1					998.47	1337.46	1395.62		2739.44
Q 8WXX5	Q8WXX5	DNJC9_HUMAN	DNAJC9	5398.79	7631.21	5760.53	6415.18	9621.47	5376.32	4944.47	4532.65	4356.71
Q 8WY22	Q8WY22	BRI3B_HUMAN	BRI3BP								832.352	
Q 8WYL5	Q8WYL5	SSH1_HUMAN	SSH1	830.693	1272.34	864.219		2872.52	1945.42	1205.28		
Q 8WZ42	Q8WZ42	TITIN_HUMAN	TTN		16565.2	4446.17	8201.53		2094.95		15912	1920.69
Q 8WZA9	Q8WZA9	IRGQ_HUMAN	IRGQ					3748.98	4467.61	1060.03		1382.43
Q 92466	Q92466	DDB2_HUMAN	DDB2		12736.7		4863.72	11121.1	5457.53	1621.17	2978.07	2594.7
Q 92485	Q92485	ASM3B_HUMAN	SMPDL3B				1898.52	2313.05	1518.1	4942.01	1514.77	6325.95
Q 92499	Q92499	DDX1_HUMAN	DDX1			3546.43	7269.99	24150.4	8691.63	3650.82	4692.45	4205.58
Q 92504	Q92504	S39A7_HUMAN	SLC39A7	10486.5	10086.1	13029	3807.62	5553.28	3874.74	6960.28	4055.07	6649.25
Q 92506	Q92506	DHB8_HUMAN	HSD17B8				1240.26			1263.11		
Q 92522	Q92522	H1X_HUMAN	H1-10	3338.34	1658.55	2306.15	1118.78	1689.54		2226.37	2631.36	2622.38
Q 92526	Q92526	TCPW_HUMAN	CCT6B	11980.3	4856.98	13328	25708.8	12447.8	30969.2	14664.6		27238.1
Q 92530	Q92530	PSMF1_HUMAN	PSMF1			2250.83	1917.26	2180.47	3948.72	9007.93	2946.26	7527.94
Q 92536	Q92536	YLAT2_HUMAN	SLC7A6							1513.16	338.252	1063.17
Q 92538	Q92538	GBF1_HUMAN	GBF1	3404.92	3515.49	3400.25	2033.79	1947.84	1797.78	2863.51	2195.24	2839.69
Q 92541	Q92541	RTF1_HUMAN	RTF1					183.016	357.406	260.344		264.723
Q 92542	Q92542	NICA_HUMAN	NCSTN	6652.56		8927.05	7771.13	6314.81	6937.88	3121.25	4681.99	2533.42
Q 92546	Q92546	RGP1_HUMAN	RGP1		511.755			951.563	1001.73	446.562		471.987
Q 92552	Q92552	RT27_HUMAN	MRPS27	4604.22	4940.33	4541.9	3223.67	7855.7	6717.83	4736.96	4628.83	3904.8
Q 92575	Q92575	UBXN4_HUMAN	UBXN4			2257.83	4339.96	8929.98	1326.77	2896.2	4103.75	2499.22
Q 92597	Q92597	NDRG1_HUMAN	NDRG1	14168.8	19147.4	21000.9	16936.3		18409.8			
Q 92598	Q92598	HS105_HUMAN	HSPH1	3859.07		6519.38	11328.9	10114.2	20019.9	10441	6881.12	11548.9
Q 92599	Q92599	SEPT8_HUMAN	SEPTIN8				2416.34	1594.33	2544.43	990.06		1087.54
Q 92600	Q92600	CNOT9_HUMAN	CNOT9				1267.38	2699.63	1636.36	2616.35	3109.9	2385.37
Q 92609	Q92609	TBCD5_HUMAN	TBC1D5	4769.93	2244	4188.4	2994.04			2517.55	3857.83	1666.23
Q 92614	Q92614	MY18A_HUMAN	MYO18A	4185.06			8307.88	8855.12	5579.48	10018.8	7586.3	7680.56
Q 92615	Q92615	LAR4B_HUMAN	LARP4B	5634.68	4290.88	5745.74	5827.94	15356.9	7584.83	10847	10711.5	11882.7
Q 92616	Q92616	GCN1_HUMAN	GCN1	24396.4	23074	18942.4	43201.9	30879.2	27774	33097	39301.4	33826

Q 96N66	Q96N66	MBOA7_HUMAN	MBOAT7		1960.2	981.151	764.019	2476.1	3051.11	1186.12		
Q 96N67	Q96N67	DOCK7_HUMAN	DOCK7	60355.7	72207.9	56375.4	67437	47107.6	31063.5	51340.9	79200.4	61040.4
Q 96NE9	Q96NE9	FRMD6_HUMAN	FRMD6	1967.73	2520.69	2062.95	806.001					519.66
Q 96P11	Q96P11	NSUN5_HUMAN	NSUN5	4409.16	9153.29	4839.51	1940.12	6319.55	2042.49	5086.38	5612.75	5691.05
Q 96P47	Q96P47	AGAP3_HUMAN	AGAP3					639.802		634.361	1065.94	612.912
Q 96P48	Q96P48	ARAP1_HUMAN	ARAP1	1335.91		2973.94	3380.03	3914.17	6555.91	3285.41	2894.56	4876.34
Q 96P63	Q96P63	SPB12_HUMAN	SERPINB12	32233.9	21268.5	54196.6	31013.6	4098.43	11224.7	9828.36	33833.6	23105.1
Q 96P70	Q96P70	IPO9_HUMAN	IPO9	4642.3	6598.72	5559.91	6380.68	7699.23	6478.58	6659.07	6432.52	8731.11
Q 96PC5	Q96PC5	MIA2_HUMAN	MIA2				1304.77	6089.17	4722.6	3954.6	1198.14	2111.22
Q 96PK6	Q96PK6	RBM14_HUMAN	RBM14	22408.9	68154.8	19781.4	22370.8	87200.6	12561.3	11760.3	18535.6	15619.2
Q 96PU5	Q96PU5	NED4L_HUMAN	NEDD4L				1113.58	490.253	958.133	1304.24	523.623	999.82
Q 96PU8	Q96PU8	QKI_HUMAN	QKI	5165.01	3653.28	4957.68	6086.99	8120.1	5186.34	1761.51	1154.44	2197.6
Q 96PV6	Q96PV6	LENG8_HUMAN	LENG8	6543.76	16012	3387.11	2074.52	11230.7	1680.07	1045.07		1213.2
Q 96PY5	Q96PY5	FMNL2_HUMAN	FMNL2	4799.79	7821.87	8531.33	10743.7	44912.5	25226.4	45611.5	8577.51	39419.5
Q 96Q11	Q96Q11	TRNT1_HUMAN	TRNT1				1590.76		1782.72	2627.37	1763.41	1612
Q 96QA5	Q96QA5	GSDMA_HUMAN	GSDMA	15070.5	3092.39	11948.6	5601.01	37940.3	5409.42	3942.35	4133.1	5615.96
Q 96QC0	Q96QC0	PP1RA_HUMAN	PPP1R10				449.87	1329.93	619.262	1371.75	270.409	886.532
Q 96QD8	Q96QD8	S38A2_HUMAN	SLC38A2	2343.91	5900.28	4058.74	1052.51	3332.29	2234.52	2963.66		3715.71
Q 96QK1	Q96QK1	VPS35_HUMAN	VPS35	3713.72	6844.26	4600.07	2444.71	3906.01	9814.73	3300.15	1967.6	3370.23
Q 96QR8	Q9UJV8;Q96QR8	PURB_HUMAN	PURB	37680.1	78644.6	22843.7	29885.3	28451.6	13402	15865.2	17127.7	19892.8
Q 96QU8	Q96QU8	XPO6_HUMAN	XPO6	4569.44	3674.72	4158.51	3111.56	1895.01	3466.85	5131.84	5190.7	4198.11
Q 96RF0	Q96RF0	SNX18_HUMAN	SNX18	5601.62	3284.86	4363.06		5270.36	4186.28			
Q 96RN5	Q96RN5	MED15_HUMAN	MED15					1825.43	1409.06	2158.06		2383.85
Q 96RP9	Q96RP9	EFGM_HUMAN	GFM1			2390.01	2972.41	3766.91	3566.72	2659.12	2335.98	3217.95
Q 96RQ3	Q96RQ3	MCCA_HUMAN	MCCC1	850938	803036	819732	591547	205670	159985	257288	719283	296687
Q 96RQ9	Q96RQ9	OXLA_HUMAN	IL4I1					1583.1	1177.51			
Q 96RS0	Q96RS0	TGS1_HUMAN	TGS1				2332.67			2401.82	4821.74	1942.02
Q 96RT1	Q96RT1	ERBIN_HUMAN	ERBIN				4373.14	9956.78	7108.45	8446.89	2133.91	9935.66
Q 96RT7	Q96RT7	GCP6_HUMAN	TUBGCP6			400.364	1343.83	2235.83	1405.23	1508.86	990.838	1286.95
Q 96RT8	Q96RT8	GCP5_HUMAN	TUBGCP5					352.231	635.045	788.172		799.688
Q 96S52	Q96S52	PIGS_HUMAN	PIGS	2837.92	2709	4088.78	1791.99	4784.98	3494.54	2972.74	1852.75	4390.84
Q 96S55	Q96S55	WRIP1_HUMAN	WRNIP1		2996.57	1341.14	3882.96	4958.8	4989.78	6439.92	5778.71	6363.23
Q 96S59	Q96S59	RANB9_HUMAN	RANBP9					1095.58	1993.37	1818.03		1623.42
Q 96S97	Q96S97	MYADM_HUMAN	MYADM	3899.46	2276.56	6420.69	14251.5	2783.76	5047.31	10647.2	9108.2	13584.1
Q 96SB4	Q96SB4	SRPK1_HUMAN	SRPK1				2276.84	2842.14	1796.01	2170	2978.01	2262.53
Q 96SI9	Q96SI9	STRBP_HUMAN	STRBP		2692.61			6325	4033.7	5662.6	2626.9	
Q 96SL4	Q96SL4	GPX7_HUMAN	GPX7									
Q 96SN8	Q96SN8	CK5P2_HUMAN	CDK5RAP2	6521.95	7718.69	3012.38		3718.78	2724.16	1370.19		1761.25
Q 96SQ9	Q96SQ9	CP2S1_HUMAN	CYP2S1				2363.15	4713.75	5077.98	7106.18	3622.87	5361.41
Q 96ST3	Q96ST3	SIN3A_HUMAN	SIN3A			1070.53	970.522	1297.65	1190.98	1179.69		838.51
Q 96SU4	Q96SU4	OSBL9_HUMAN	OSBPL9				1774.61	8669.28	11481.2	3544.79		
Q 96SZ6	Q96SZ6	CK5P1_HUMAN	CDK5RAP1	2273.31			2775.41	8122.01	4131.87	7559.61	7152.53	7354.45
Q 96T23	Q96T23	RSF1_HUMAN	RSF1				869.74	2665.49	1201.32	173.498		592.202
Q 96T51	Q96T51	RUFY1_HUMAN	RUFY1				760.635	2227.07	1575.17	1705.44		
Q 96T59	Q96T59	CDRTF_HUMAN	CDRT15	5742.8	7933.92	4046.5	4478.85	23724.7	11418.5	7908.38	2687.36	7174.82
Q 96T60	Q96T60	PNKP_HUMAN	PNKP	5122.73	5166		6838.48	22852.5	3179.88	1797.85	762.659	847.297
Q 96T76	Q96T76	MMS19_HUMAN	MMS19	2216.95	3388.2	2607.57	4284.24	6022.61	7254.11	2882.63	2937.9	3031.4
Q 96T88	Q96T88	UHRF1_HUMAN	UHRF1				3350.56	2639.53	1756.21	2242.44	2816.67	2686.23
Q 96TA1	Q96TA1	NIBA2_HUMAN	NIBAN2	23950.5	7246.5	10302.4	7425.07	9740.92	10037	4479.65	4937.08	4980.67
Q 96TA2	Q96TA2	YMEL1_HUMAN	YME1L1					3833.95			5221.57	
Q 96TC7	Q96TC7	RMD3_HUMAN	RMDN3							30173.1		41172.4
Q 99081	Q99081	HTF4_HUMAN	TCF12					1798.77				
Q 99436	Q99436	PSB7_HUMAN	PSMB7	2436.18		2194.95	1137.59	2840.01	8302.03	3145.69	533.658	3702.11
Q 99439	Q99439	CNN2_HUMAN	CNN2	11095.4	9216.61	9429.83	15702.7	10457.2	5119.97	5438.44	8198.34	5423.41
Q 99442	Q99442	SEC62_HUMAN	SEC62	2383.02			1399.17	1941.47	2420.71	3884.5	1339.74	3450.89
Q 99459	Q99459	CDC5L_HUMAN	CDC5L			2082.44	3094	6291.36	1984.26	2767.82	1233.34	3514.61
Q 99460	Q99460	PSMD1_HUMAN	PSMD1	7957.35	6863.23	10238	12725.4	13304.8	15873.6	11152.5	9609.83	11315.2
Q 99470	Q99470	SDF2_HUMAN	SDF2					3457.88	3549.78	368.437		
Q 99471	Q99471	PFD5_HUMAN	PFDN5						4733.93	790.339		1263.42
Q 99497	Q99497	PARK7_HUMAN	PARK7	3787.62		4265.59	1900.19	1738.5	5901.65	2291.06	2254.32	2498.64
Q 99536	Q99536	VAT1_HUMAN	VAT1	16412.5	7960.63	19267.3	15222.7	12324	31401.1	9510.8	7382.68	7366.9
Q 99538	Q99538	LGMN_HUMAN	LGMN					1406.27	2594.89	822.163		2255.11
Q 99543	Q99543	DNJC2_HUMAN	DNAJC2					3189.6	4254.33	2416.94		2556.42
Q 99567	Q99567	NUP88_HUMAN	NUP88				3167.97	5647.43	5462.97	6824.45	1744.09	8387.65
Q 99569	Q99569	PKP4_HUMAN	PKP4	1523.48				2859.54	1833.67	3810.48	738.186	3759.15
Q 99570	Q99570	PI3R4_HUMAN	PIK3R4				5307.81	2863.13	3495.29	4824.77	4595.99	3926.52
Q 99575	Q99575	POP1_HUMAN	POP1		2699.51		2988.67	5864.94	4774.67	4823.35	2876.79	4431.47
Q 99584	Q99584	S10AD_HUMAN	S100A13	5483.38		3595.79	4470.73	2197.9	6507.04	3370.2	3796.76	2928.88
Q 99615	Q99615	DNJC7_HUMAN	DNAJC7				2673.12	4894.55	4232.41	2568.64		3652.01
Q 99623	Q99623	PHB2_HUMAN	PHB2	6875.7	5564.45	6017.43	4754.46	14133.8	9044.59	15466.7	8090.24	15174.4
Q 99627	Q99627	CSN8_HUMAN	COPS8			4467.82	2339.65	1562.48	3026.48	2600.4		2068.79
Q 99643	Q99643	C560_HUMAN	SDHC		1829.09					2454.22	1043.56	1431.9
Q 99653	Q99653	CHP1_HUMAN	CHP1	5437.95	6392.7	6294.97	6696.72	6130.31	7598.35	7472.52	7199.33	6412.67
Q 99661	Q99661	KIF2C_HUMAN	KIF2C				7628.47		3531.38	6197.02	5293.88	5224.56
Q 99700	Q99700	ATX2_HUMAN	ATXN2	7967.14	6845.61		3766.18	12529.2	3366.87	2039.44	2904.77	2364.65
Q 99707	Q99707	METH_HUMAN	MTR				6401.53		2224.08	2948.99	5603.25	2940.33
Q 99714	Q99714	HCD2_HUMAN	HSD17B10	5105.43	1891.96	5905.61	3691.2		5245.53	6550.72	6835.79	12136.3
Q 99729	Q99729	ROAA_HUMAN	HNRNPAB	7068.87	3988.91	1724.39	4768.82	11739	8564.16	5244.75	2717.64	5995.57
Q 99733	Q99733	NP1L4_HUMAN	NAP1L4	6230.44		5682.82	5531.95	3124.74	7718.92	5459.89	2405.61	6770.94
Q 99747	Q99747	SNAG_HUMAN	NAPG			1631.55	1701.97	844.737	919.84	1070.14		614.379
Q 99755	Q99755	PI51A_HUMAN	PIP5K1A	4346.09	5667.06	3157.03	3047.05	8312.1	3870.82	1438.86		1450.91
Q 99798	Q99798	ACON_HUMAN	ACO2			3741.34	3279.6	4803.78	6129.44	2924.44	2465.32	2885.39
Q 99811	P54821;Q99811	PRRX2_HUMAN	PRRX2									
Q 99816	Q99816	TS101_HUMAN	TSG101	8705		8139.34	13760.6	11173.5	14224.1	20377.3	6760.79	16788.8
Q 99829	Q99829	CPNE1_HUMAN	CPNE1	5273.01	3510.87	3800.94	2057.97	6896.3	12866.4	6223.53	2822.68	4813.31
Q 99832	Q99832	TCPH_HUMAN	CCT7	8811.4	7717.15	9165.48	15084.9	15746.2	21973.6	25790.2	12581.2	22465.5
Q 99836	Q99836	MYD88_HUMAN	MYD88	7805.93	8683.95	7012.08	5575.1		6103.01	5324.53	4034.25	5821
Q 99848	Q99848	EBP2_HUMAN	EBNA1BP2		1848.56		1028.56	2539.95	1108.33	1454		989.541
Q 99856	Q99856	ARI3A_HUMAN	ARID3A				1050.97	2047.12	956.389			
Q 99873	Q99873	ANM1_HUMAN	PRMT1	2587.57		2063.76	2566.49	3550.91	6449.63	3365.27	1747.29	4075.97
Q 99958	Q99958	FOXC2_HUMAN	FOXC2									
Q 99959	Q99959	PKP2_HUMAN	PKP2				2068.98	7287.06	3326.41	6571	1888.91	7264.18
Q 9BPW8	Q9BPW8	NIPS1_HUMAN	NIPSNAP1		7271.59	8555.17	4617.79	5748.5	4745.66	3683.09		4618.1
Q 9BPX3	Q9BPX3	CND3_HUMAN	NCAPG	2963.72	2539.23	2744.84	2170.69	5329.88	4679.89	3229.79	4529.5	3406.5
Q 9BPX5	Q9BPX5	ARP5L_HUMAN	ARPC5L	6805.97		7353.54	7395.76	6828.63	2554.04	5754.74	7039.79	5319.8
Q 9BPX6	Q9BPX6	MICU1_HUMAN	MICU1							1419.06		1537.35
Q 9BQ13	Q9BQ13	KCD14_HUMAN	KCTD14							1767.76	2316.22	2200.99
Q 9BQ39	Q9BQ39	DDX50_HUMAN	DDX50		2511.78	2049.95		3999.05	1195.33	3956.01	2630.17	4989.38
Q 9BQ67	Q9BQ67	GRWD1_HUMAN	GRWD1				4746.99	4999.51	1913.84	1415.78	3408.16	2111.17
Q 9BQ69	Q9BQ69	MACD1_HUMAN	MACROD1							3477.61		3821.09
Q 9BQ70	Q9BQ70	TCF25_HUMAN	TCF25	1511.76	1409.57	1070.01				861.034		1170.38
Q 9BQA1	Q9BQA1	MEP50_HUMAN	WDR77				1028.52		1633.02	952.864	2884.25	
Q 9BQC3	Q9BQC3	DPH2_HUMAN	DPH2				1817.54	1841.06	2783.85	1829.77	882.981	1789.42
Q 9BQE3	P68366;P68363;Q9BQE3;Q71U36;A6NHL2	TBA1C_HUMAN	TUBA1C	648857	578516	627288	799376	527215	476319	788648	937399	927101
Q 9BQE5	Q9BQE5	APOL2_HUMAN	APOL2				7941.52	10187.8	7125.03	2713.52	2106.45	3270.19
Q 9BQG0	Q9BQG0	MBB1A_HUMAN	MYBBP1A	1976.75	1763.82		753.205	3295.92	3227.56	4772.28	4779.22	3045.27
Q 9BQP7	Q9BQP7	MGME1_HUMAN	MGME1	882.886	2643.56	3108.37	1919.25	2426.29	4835.12	2631.7	1627.7	3261.84
Q 9BR76	Q9BR76	COR1B_HUMAN	CORO1B	7221.5	6873.05	7648.6	11685.1	51624.9	6930.92	12615.8	12330.5	6669.33
Q 9BRA2	Q9BRA2	TXD17_HUMAN	TXNDC17	2097.28		1146	1532.29	799.566	2862.54	2491.28	1518.05	1904.44
Q 9BRG1	Q9BRG1	VPS25_HUMAN	VPS25				1077.3	913.23	1604.05	1337.46	1013.98	1782.28
Q 9BRJ6	Q9BRJ6	CG050_HUMAN	C7orf50	1943.78	2121.88	1895.34	1652.87	4292.69	4791.68	5017.79	3908.46	5913.06
Q 9BRK4	Q9BRK4;O60513	LZTS2_HUMAN	LZTS2	6618.76	7639.16	6706.95	2252.49	3004.98	2261.88	3210.93		2315.05
Q 9BRK5	Q9BRK5	CAB45_HUMAN	SDF4			3729.86	2924.82		1905.38	4141.44	5090.64	5462.53
Q 9BRP4	Q9BRP4	PAAF1_HUMAN	PAAF1		2473.68		5563.8	5451.99	10457	7292.58	3557.52	7141.32
Q 9BRP8	Q9BRP8	PYM1_HUMAN	PYM1	2357.47		2955.25	3496.67	3247.3	4851.21	4909.22	2885.22	3935.78
Q 9BRQ6	Q9BRQ6	MIC25_HUMAN	CHCHD6	2650.65	4190.59	3307.84			459.984			597.14
Q 9BRQ8	Q9BRQ8	FSP1_HUMAN	AIFM2				2479.53		1479.42	5738.75	2250.86	4697.93
Q 9BRR6	Q9BRR6	ADPGK_HUMAN	ADPGK					1560.33	1071.02			436.493
Q 9BRT6	Q9BRT6	LLPH_HUMAN	LLPH	3958.52	6372.67	3551.24	1791.4	14289.6	6592.53	2054.74	1532.82	1622.34
Q 9BRX2	Q9BRX2	PELO_HUMAN	PELO		2648.58			1825.03	2042.01	1607.41	2310.91	2349.25
Q 9BS26	Q9BS26	ERP44_HUMAN	ERP44					2288.31	3107.62	1740.62		1005.5
Q 9BSD7	Q9BSD7	NTPCR_HUMAN	NTPCR	10319.3	7716.34	10698	10946.2	7404.81	5637.4	5017.13	6219.72	5686.12
Q 9BSH4	Q9BSH4	TACO1_HUMAN	TACO1	983.421		1498.94	581.463	1864.12	2638.54	1145.52		785.565
Q 9BSJ2	Q9BSJ2	GCP2_HUMAN	TUBGCP2	3068.52	2544.05	4006.06	4317.26	4398.79	4711.74	6429.12	4810.52	6860.55
Q 9BSJ8	Q9BSJ8	ESYT1_HUMAN	ESYT1	4752.6		6491.51	10205.7	7084.5	8441.97	16064.8	10609.5	17378.2
Q 9BST9	Q9BST9	RTKN_HUMAN	RTKN							2489.9	2906.23	3728.4
Q 9BSV6	Q9BSV6	SEN34_HUMAN	TSEN34					1242.11	586.076	789.632		
Q 9BT22	Q9BT22	ALG1_HUMAN	ALG1	7932.31	9849.89	11040.8	5576.08	4894.28	5945.16	10409.2	11564.4	12459.4
Q 9BT40	Q9BT40	INP5K_HUMAN	INPP5K				1004.97	1406.08	1197.42	2343.57		2554.1
Q 9BT78	Q9BT78	CSN4_HUMAN	COPS4	4457.15		7423.64	5814.56	3713.16	5016.62	5998.17	4170.56	5677.59
Q 9BTD8	Q9BTD8	RBM42_HUMAN	RBM42	2108.78				2895.37	2255.43	1070.64		1495.48
Q 9BTE3	Q9BTE3	MCMBP_HUMAN	MCMBP				962.859	1562.24	2290.23	1632.18	850.195	1985.07
Q 9BTE6	Q9BTE6	AASD1_HUMAN	AARSD1						2177.55			756.741
Q 9BTT0	Q9BTT0	AN32E_HUMAN	ANP32E			3807.84	3113.05		4205.43	2570.74	1374.78	2890.35
Q 9BTT6	Q9BTT6	LRRC1_HUMAN	LRRC1					5151.18	3999.02	7309.9	1764.7	4143.79
Q 9BTU6	Q9BTU6	P4K2A_HUMAN	PI4K2A	3412.43		3545.29	3461.15	5435.54	3983.37	5521.09	3188.27	4071.91
Q 9BTV4	Q9BTV4	TMM43_HUMAN	TMEM43	3648.37	3749.98	4525.75	2726.25	2879.18	2766.03	3139.94	2210.63	4390.26
Q 9BTW9	Q9BTW9	TBCD_HUMAN	TBCD	6216.7	4037.52	5169.83	8397.38	4445.15	6168	4929.48	11536	6352.03
Q 9BTX1	Q9BTX1	NDC1_HUMAN	NDC1				1453.18	801.742	539.807	1595.94	962.749	1080.42
Q 9BTY7	Q9BTY7	HGH1_HUMAN	HGH1			1759.35	4482.15			4840.76	4567.98	5205.88
Q 9BUB7	Q9BUB7	TMM70_HUMAN	TMEM70						1516.69	2495.69	2881.9	2829.32
Q 9BUF5	Q9BUF5;Q13509	TBB6_HUMAN	TUBB6	190529	101736	181636	195293	54230.8	53393.8	49350.1	55369.8	38265.2
Q 9BUH6	Q9BUH6	PAXX_HUMAN	PAXX	8831.32	5804.32		8068.07	6177.66	11517.6	3872.37	3727.19	2078.73
Q 9BUI4	Q9BUI4	RPC3_HUMAN	POLR3C	5574.36	5482.88		650.895	1195.05	739.598	604.956		467.12
Q 9BUJ2	Q9BUJ2	HNRL1_HUMAN	HNRNPUL1	15471.7	21728.3	18266.9	5353.39	28665.7	10700.4	9749.53	7823.64	9583.48
Q 9BUK6	Q9BUK6	MSTO1_HUMAN	MSTO1				1388.46	5058.85	1304.62	2085.37	965.27	2118.1
Q 9BUL8	Q9BUL8	PDC10_HUMAN	PDCD10				2905.7	2133.59	2561.72	1858.32	1240.78	1144.58
Q 9BUN8	Q9BUN8	DERL1_HUMAN	DERL1		5471.27	2677.43	3041.73			3936.5	1453.34	4092.59
Q 9BUQ8	Q9BUQ8	DDX23_HUMAN	DDX23		2474.75		1466.48	2738.3	1769.85	7014.64	4172.95	6370
Q 9BV40	Q9BV40	VAMP8_HUMAN	VAMP8				3880.06	3982.59	3894.51	7350.31		6342.94
Q 9BV44	Q9BV44	THUM3_HUMAN	THUMPD3	4046.41			3943.6			1752.73	3057.36	
Q 9BV68	Q9BV68	RN126_HUMAN	RNF126				3461.35			5011.88	4885.28	5878.3
Q 9BV86	Q9BV86	NTM1A_HUMAN	NTMT1					2041.75	3520.59	1215.84	1097.89	
Q 9BVC4	Q9BVC4	LST8_HUMAN	MLST8					644.558	1123.02	2293.92	1261.93	1611.44
Q 9BVG4	Q9BVG4	PBDC1_HUMAN	PBDC1			14645	6520.53	5301.78	8193.28	3870.01	4816.87	4309.95
Q 9BVG8	Q9BVG8	KIFC3_HUMAN	KIFC3		1433.32			2225.76	713.292	1931.13		2228.24
Q 9BVI4	Q9BVI4	NOC4L_HUMAN	NOC4L	2159.2			1934.14	2696.18	2025.46	4247.61	2853.41	3978.41
Q 9BVK6	Q9BVK6	TMED9_HUMAN	TMED9			9743.97	134818	10938.1	13020.4	20596.5	4108.51	17231.7
Q 9BVL2	Q9BVL2	NUP58_HUMAN	NUP58				438.193	1317.81	2110.07	1016.27		
Q 9BVP2	Q9BVP2	GNL3_HUMAN	GNL3	5805.41			4595.87		4814.48	7416.7	4292.15	7102.69
Q 9BVQ7	Q9BVQ7	SPA5L_HUMAN	SPATA5L1	4845.45	4868.37	3822.77	5453.01	7676.89	3417.38	4766.49	3037.84	3729.32
Q 9BVS4	Q9BVS4	RIOK2_HUMAN	RIOK2					2440.37	834.86			
Q 9BW04	Q9BW04	SARG_HUMAN	SARG								779.511	528.99
Q 9BW19	Q9BW19	KIFC1_HUMAN	KIFC1				2634.72	2484.34	1813.38	2017.56	2798.04	2021.13
Q 9BW83	Q9BW83	IFT27_HUMAN	IFT27	3162.17	1396.16	3822.75				1651.82		1841.19
Q 9BW92	Q9BW92	SYTM_HUMAN	TARS2	3702.14		3538.01	7833.44	5646.13	5034.06	8732	9096.81	9522.18
Q 9BWF3	Q9BWF3;Q9BQ04	RBM4_HUMAN	RBM4	6431.29	6167.04	8212.92	4887.02			3527.93	6834.03	1861.55
Q 9BWH6	Q9BWH6	RPAP1_HUMAN	RPAP1			3087.29	6178.46	4696.1	5644.8	5862.02	4247.2	5309.03
Q 9BWJ5	Q9BWJ5	SF3B5_HUMAN	SF3B5				1480.64	797.148	668.462	689.635	1484.77	
Q 9BWS9	Q9BWS9	CHID1_HUMAN	CHID1					968.533	3207.64	2677.49		
Q 9BX10	Q9BX10	GTPB2_HUMAN	GTPBP2				2745.65	6374.2	3544.65	2312.67		3410.26
Q 9BX40	Q9BX40	LS14B_HUMAN	LSM14B		3980.89	3543.03	2351.24		1918.09	3306.78		
Q 9BX46	Q9H0Z9;Q9BX46	RBM24_HUMAN	RBM24				1449.69	5012.38	1701.34			
Q 9BX68	Q9BX68	HINT2_HUMAN	HINT2			1481.25			2389.33	522.058		451.772
Q 9BXB5	Q9BXB4;Q9BXB5	OSB10_HUMAN	OSBPL10		868.657		3338.78	3317.53	3872.35	3821.84	2206.4	2272.97
Q 9BXJ9	Q9BXJ9	NAA15_HUMAN	NAA15	5671.35		3778.53	6287.67	5850.12	6571.14	4142.79		3479.28
Q 9BXP2	Q9BXP2	S12A9_HUMAN	SLC12A9					5955.17	4073.37	1813.87		
Q 9BXP5	Q9BXP5	SRRT_HUMAN	SRRT			1300.25		6921.17	5094.85	4840.93		3073.7
Q 9BXS5	Q9BXS5	AP1M1_HUMAN	AP1M1	9452.04	16482.8	9862.65	8823.87	7972.85	10771.4	11983	6390.07	12004.7
Q 9BXS6	Q9BXS6	NUSAP_HUMAN	NUSAP1							776.279	1503.49	856.332
Q 9BXW6	Q9BXW6	OSBL1_HUMAN	OSBPL1A	10907	2958.37	13398.5	25714		11956.5	9864.76	24485.3	19842.8
Q 9BXW7	Q9BXW7	HDHD5_HUMAN	HDHD5			7679.31	2000.61		2563.47	3675.12	1792.16	2592.22
Q 9BXW9	Q9BXW9	FACD2_HUMAN	FANCD2				390.33	2287.95	931.943	1651.18	1170.29	2378.45
Q 9BY44	Q9BY44	EIF2A_HUMAN	EIF2A	3827.58		3369.54	5974.98	3107.66	3684.34	2856.65	4080.63	2987
Q 9BY77	Q9BY77	PDIP3_HUMAN	POLDIP3	8610.14	8361.62	11022.1	5728.82	7573.59	5960.98	3776.34	11982.2	6370.12
Q 9BY89	Q9BY89	K1671_HUMAN	KIAA1671					694.612		813.331		
Q 9BYD2	Q9BYD2	RM09_HUMAN	MRPL9	1572.36	1185.81	1664.94	1130.75	491.664	618.816	1544.24	1109.14	1644.46
Q 9BYD3	Q9BYD3	RM04_HUMAN	MRPL4						2941.68	4332.67	2094.25	4020.52
Q 9BYI3	Q9BYI3	HYCCI_HUMAN	HYCC1					6084.85	3320.37	1767.24		2687.24
Q 9BYJ1	Q9BYJ1	LOXE3_HUMAN	ALOXE3	9896.78	12536.8	14728.5	4085.36		5506.36	4043.54	4338.17	6833.81
Q 9BYK8	Q9BYK8	HELZ2_HUMAN	HELZ2					3901.17	1716			
Q 9BYX7	Q6S8J3;P0CG38;Q9BYX7;P0CG39	ACTBM_HUMAN	POTEKP	22280.2	13642	28117.5	16099.4	24188.9	20609	26599	4392.29	31545.2
Q 9BZE1	Q9BZE1	RM37_HUMAN	MRPL37	5066.76				3987.89	5233.41	6010.52	4428.04	4442.29
Q 9BZE4	Q9BZE4	GTPB4_HUMAN	GTPBP4	8717.56		7190.19	6346.71		7238.54	6551.41	8044.48	5834.57
Q 9BZF1	Q9BZF1	OSBL8_HUMAN	OSBPL8		2658.93	2028.3	1335.31	2999.18	3278.22	6371.59	2130.36	5948.94
Q 9BZF9	Q9BZF9	UACA_HUMAN	UACA	3246.33	11008	8564.3	46350	100091	53793.3	8111.59	6194.36	6538.96
Q 9BZH6	Q9BZH6	WDR11_HUMAN	WDR11					1645.23	1293.5	1237.87		1673.48
Q 9BZI7	Q9BZI7	REN3B_HUMAN	UPF3B					928.003	571.784	1050.12	790.422	
Q 9BZJ0	Q9BZJ0	CRNL1_HUMAN	CRNKL1				3318.79				5154.9	
Q 9BZL4	Q9BZL4	PP12C_HUMAN	PPP1R12C				778.882	3164.04	1535.59			
Q 9BZL6	Q9BZL6	KPCD2_HUMAN	PRKD2	976.106	1260.71	1421.4	2592.43	3644.98	4341.78	5508.9	1952.69	4892.96
Q 9BZM5	Q6H3X3;Q9BZM5;Q5VY80	ULBP2_HUMAN	ULBP2					2360.76		5519.27	5957.62	5116.49
Q 9BZQ8	Q9BZQ8	NIBA1_HUMAN	NIBAN1	5006.28		6579.35	3342.22	1615.48	2071.96			
Q 9BZX2	Q9BZX2	UCK2_HUMAN	UCK2		4771.48	2813.51	9137.19	6522.9	9705.06	13462.8	14115.9	14238.2
Q 9BZZ5	Q9BZZ5	API5_HUMAN	API5	6803.1		5945.01	6390.39	2916.2	6961.35	6813.39	4667.48	7867.49
Q 9C005	Q9C005	DPY30_HUMAN	DPY30				8525.03	7951.42	3951.27	4521.08	6993.55	4599.4
Q 9C037	Q9C037	TRIM4_HUMAN	TRIM4	3285.53	3755.42	3779.53	1895.53			2608.55	2424.72	2188.81
Q 9C075	Q9C075	K1C23_HUMAN	KRT23			7503.62	4805.82		2464.37	14980.2	7233.85	15915.7
Q 9C0B5	Q9C0B5	ZDHC5_HUMAN	ZDHHC5			2276.95	3385.86	5737.44	4147.1	13184.5	4469.6	11538.3
Q 9C0B7	Q9C0B7	TNG6_HUMAN	TANGO6				3727.3	3092.01	2251.79	4644.88	8887.48	3541.03
Q 9C0C2	Q9C0C2	TB182_HUMAN	TNKS1BP1	15098.6	23181.4	12913	7753.52	11450.8	6783.78	4804.21	6377.55	6376.43
Q 9C0C9	Q9C0C9	UBE2O_HUMAN	UBE2O				3026.45	4182.76	3914.32	1612.76		3459.38
Q 9C0E2	Q9C0E2	XPO4_HUMAN	XPO4				3133.66	3495.14		2484.32	3010.76	2641.98
Q 9GZL7	Q9GZL7	WDR12_HUMAN	WDR12						1296.14	1252.59		1500.26
Q 9GZS3	Q9GZS3	SKI8_HUMAN	SKIC8				5334.35	2886	5380.93	6930.14	5827.39	6599.75
Q 9GZT3	Q9GZT3	SLIRP_HUMAN	SLIRP	6098.19	5020.98	5109.44	8008.28	6574.48	6486.8	13591.8	17903.5	9265.68
Q 9GZT8	Q9GZT8	NIF3L_HUMAN	NIF3L1					2398.54	3746.07	2021.15		1555.98
Q 9GZY6	Q9GZY6	NTAL_HUMAN	LAT2				1293.3	2031.01	1360.33	4111.5	1077.82	5449.21
Q 9GZZ1	Q9GZZ1	NAA50_HUMAN	NAA50	5142.55		5858.64	1821.23	3339.94	7063.22	3693.14		2438.27
Q 9GZZ8	Q9GZZ8	LACRT_HUMAN	LACRT								13812.4	
Q 9GZZ9	Q9GZZ9	UBA5_HUMAN	UBA5				6619.37	9583.54	3283.22	4556.89	6864.67	4645.45
Q 9H019	Q9H019	MFR1L_HUMAN	MTFR1L	8604.44	6065.28	6108.55	2938.89	3093.73	2881.7	2764.97	1620.98	3281.15
Q 9H0A0	Q9H0A0	NAT10_HUMAN	NAT10	1480.55		1924.16	1064.45	530.406	642.363	2969.49	3068.59	3509.32
Q 9H0D6	Q9H0D6	XRN2_HUMAN	XRN2	25140.1	38351	7767.3	8652.65	15682.1	8508.84	9957.59	10654.2	10658.2
Q 9H0E2	Q9H0E2	TOLIP_HUMAN	TOLLIP					3260.52	3321.66	3712.64		1501.17
Q 9H0H5	Q9H0H5	RGAP1_HUMAN	RACGAP1		2946.15	3252.62		7337.93	3731.31	1278.47	1307.43	1714.6
Q 9H0K1	Q9H0K1	SIK2_HUMAN	SIK2					1413.17	1054.84	910.645		
Q 9H0P0	Q9H0P0	5NT3A_HUMAN	NT5C3A					2785.93	664.438	4476.4		2070.63
Q 9H0R4	Q9H0R4	HDHD2_HUMAN	HDHD2			392.491			1205.15			
Q 9H0S4	Q9H0S4	DDX47_HUMAN	DDX47	4526.97	3703.27	2256.1	4197.41	3864.5	4315.95	6161.74	6861.75	4220.66
Q 9H0U3	Q9H0U3	MAGT1_HUMAN	MAGT1				962.138		1560.43	6408.69		3071.16
Q 9H0U4	Q92928;Q9H0U4	RAB1B_HUMAN	RAB1B	36531.4	30283.9	38812	29494.2	91133.4	38770.7	71804.9	26111.4	63375
Q 9H0W8	Q9H0W8	SMG9_HUMAN	SMG9	3190.85		2490.72	3760.85	3035.34	3185.19	2300.19	3580.67	2500.59
Q 9H0W9	Q9H0W9	CK054_HUMAN	C11orf54			3507.56	2320.98	2550.44	4255.68	3010.76	2287.73	3944.56
Q 9H147	Q9H147	TDIF1_HUMAN	DNTTIP1				806.662	1358.42	1099.8	709.321		
Q 9H1A4	Q9H1A4	APC1_HUMAN	ANAPC1	1268.19			1008.95	1462.91	1097.04	1741.89	1074.83	2016.01
Q 9H1D9	Q9H1D9	RPC6_HUMAN	POLR3F				2245.07					
Q 9H1E1	Q9H1E1	RNAS7_HUMAN	RNASE7	15374.2	18664	27048.8	8460.55	4411.12	9646.46	6838.84	7817.8	13873
Q 9H1E3	Q9H1E3	NUCKS_HUMAN	NUCKS1				3527.73				3275.73	
Q 9H1K4;Q 9H936	Q9H1K4;Q9H936	GHC1_HUMAN;GHC2_HUMAN	SLC25A18;SLC25A22				1989.82			1369.06	3988.79	
Q 9H1Y0	Q9H1Y0	ATG5_HUMAN	ATG5				986.386		1678.9	1229	712.296	1138.94
Q 9H223	Q9NZN3;Q9H4M9;Q9H223	EHD4_HUMAN	EHD4	8000.38	5989.72	8916.1	15758.3	4984.1	5824.93	6000.78	10840	6305.69
Q 9H267	Q9H267	VP33B_HUMAN	VPS33B					605.024	539.629	422.425		
Q 9H269	Q9H269	VPS16_HUMAN	VPS16				2811.62	1831.6	1427.43	1174.15	1724.1	926.097
Q 9H299	Q9H299	SH3L3_HUMAN	SH3BGRL3				1976.79	2781.01	4389.68	2074.79	783.828	1498.06
Q 9H2C0	Q9H2C0	GAN_HUMAN	GAN	19101.4	11419.2	20952.8	4726.74	9728.88	5477.11	33660.6	10862.8	41923.7
Q 9H2D6	Q9H2D6	TARA_HUMAN	TRIOBP	8179.8	9035.62	6183.26	4429.66	5477.87	3758.81	1471.13		1282.64
Q 9H2G2	Q9H2G2	SLK_HUMAN	SLK	101416	69084.1	257474	74346.2	27947.8	76169.6	54947.7	76982.5	103044
Q 9H2M9	Q9H2M9	RBGPR_HUMAN	RAB3GAP2	1549.02	958.261	628.377	2295.31	2129.03	2630.45	2381.84	1667.3	2354.01
Q 9H2P0	Q9H2P0	ADNP_HUMAN	ADNP				967.856	1001.77	1179.98	2051.8	1610.59	1920.49
Q 9H2U1	Q9H2U1	DHX36_HUMAN	DHX36	2703.11	2992.87		2862.36	13420.8	4183.59	1977.36	2028.52	1930.4
Q 9H2U2	Q9H2U2	IPYR2_HUMAN	PPA2			11363.2			3677.42	7130.35	4159.76	5094.98
Q 9H2V7	Q9H2V7	SPNS1_HUMAN	SPNS1				2390.12				2114.74	
Q 9H2X9	Q9H2X9	S12A5_HUMAN	SLC12A5									
Q 9H307	Q9H307	PININ_HUMAN	PNN				2111.61	8172.96		2174.42	2006.84	
Q 9H329	Q9H329	E41LB_HUMAN	EPB41L4B							2839.65		2850.1
Q 9H3G5	Q9H3G5	CPVL_HUMAN	CPVL				6176.88	3321.93	3149.19			
Q 9H3M7	Q9H3M7	TXNIP_HUMAN	TXNIP	5464.24	8787.91	5210.3		4255.62	1706.49	36035.5	10169	22602.6
Q 9H3N1	Q9H3N1	TMX1_HUMAN	TMX1	4403.98	4726.5	5829.77	5623.89	7386.79	12470.6	6308.49	5583.91	5266.83
Q 9H3P2	Q9H3P2	NELFA_HUMAN	NELFA	3671.21	1512.46	2038.28	2055.21	2626.19	2315.79	2933.02	1608.4	3262.31
Q 9H3P7	Q9H3P7	GCP60_HUMAN	ACBD3				2541.27				1289.21	
Q 9H3R2	Q9H3R2	MUC13_HUMAN	MUC13							8716.46	5916.74	7625.91
Q 9H3U1	Q9H3U1	UN45A_HUMAN	UNC45A	10572.7	8125.41	10014.1	18395.4	15284.8	8009.4	11614.9	13447.4	10703.7
Q 9H425	Q9H425	CA198_HUMAN	C1orf198				1135.92	2095.9	1676.21	2652.35		2053.22
Q 9H444	Q9H444	CHM4B_HUMAN	CHMP4B	2673.08	2143.51		3075.93		1646.06	3101.79	3036.08	
Q 9H488	Q9H488	OFUT1_HUMAN	POFUT1				2743.03	1570.03	2562.11			
Q 9H4A3	Q9H4A3	WNK1_HUMAN	WNK1				1481.25	1185.83	1225.29	2673.55	1871.1	3389.08
Q 9H4A4	Q9H4A4	AMPB_HUMAN	RNPEP	4176.25		4052.3	3311.54		1879.93	1603.73		3028.18
Q 9H4B6	Q9H4B6	SAV1_HUMAN	SAV1			4753.12	1466.06	2729.59	2746.3	4007.47		4787.29
Q 9H4G4	Q9H4G4	GAPR1_HUMAN	GLIPR2				6620.51	4610.6	4440.76	940.161	3229.28	
Q 9H4H8	Q9H4H8	FA83D_HUMAN	FAM83D				1612.81	3323.42	2265.29	2440.84		2611.74
Q 9H4L5	Q9H4L5	OSBL3_HUMAN	OSBPL3				1650.98	2298.72	1346.9	3946.21	2029.49	3893.46
Q 9H4L7	Q9H4L7	SMRCD_HUMAN	SMARCAD1		1117.94		1549.69	3060.01	2088.41	4159.97	1972.71	3105.31
Q 9H4M9	Q9NZN3;Q9H4M9	EHD1_HUMAN	EHD1	6349.1	6980.28	7946.79	5398.72	6084.91	6002.35	4266.37	4175.26	4156.6
Q 9H4P4	Q9H4P4	RNF41_HUMAN	RNF41		5469.38		13163.7	8878.24	8493.53	5892.12	7257.99	7439.61
Q 9H553	Q9H553	ALG2_HUMAN	ALG2	11770.6	8652.95	9196.89	5973.27	5570.15	4863.32	5890.93	5679.47	6844.5
Q 9H583	Q9H583	HEAT1_HUMAN	HEATR1					5148.4	1945.06	2680.78	1494.98	4956.86
Q 9H5Q4	Q9H5Q4	TFB2M_HUMAN	TFB2M	5625.74	4065.16	4515.6	9071.61	17747.8	10909.6	9427.78	10646	9465.41
Q 9H5V8	Q9H5V8	CDCP1_HUMAN	CDCP1							21480.4	9170.53	19872.4
Q 9H5Z1	Q9H5Z1	DHX35_HUMAN	DHX35	3329.2	1980.68	2679.06	3471.01			1345.13	1467.65	1786.28
Q 9H694	Q9H694	BICC1_HUMAN	BICC1									
Q 9H6R7	Q9H6R7	WDCP_HUMAN	WDCP				1901.09		873.69	742.103	2402.67	1203.78
Q 9H6S3	Q9H6S3	ES8L2_HUMAN	EPS8L2			5564.56	7277.34		3110.38	4043.93	6321.92	4543.36
Q 9H6T3	Q9H6T3	RPAP3_HUMAN	RPAP3				2757.99		3618.36	2989.02		5045.69
Q 9H6V9	Q9H6V9	LDAH_HUMAN	LDAH				1210.19	4711.81	4137.7	5116.31		4508.63
Q 9H6W3	Q9H6W3	RIOX1_HUMAN	RIOX1					3197.94		2081.77		1761.17
Q 9H773	Q9H773	DCTP1_HUMAN	DCTPP1						591.619			
Q 9H7B4	Q9H7B4	SMYD3_HUMAN	SMYD3		2386.69	2017.24	1735.69		1882.97	2103.76	2121.95	1655.9
Q 9H7D7	Q9H7D7	WDR26_HUMAN	WDR26	5858.72	12696.8	9091.15	4511.14	4584.25	6869.71	4537.27		4643.67
Q 9H7E9	Q9H7E9	CH033_HUMAN	C8orf33				1951.38			745.133	2827.87	779.057
Q 9H814	Q9H814	PHAX_HUMAN	PHAX					6201.73	3276.49	1079.21		
Q 9H845	Q9H845	ACAD9_HUMAN	ACAD9	4858.87	5881.37	5750.54	13005.2		2289.57	4626.26	3677.63	5103.68
Q 9H857	Q9H857	NT5D2_HUMAN	NT5DC2	7576.27	5898.32	4461.61	2629.37	6238.37	3816.84	8635.29	6148.55	4799.61
Q 9H8H0	Q9H8H0	NOL11_HUMAN	NOL11				800.406	1580.54	1640.32	1770.56	784.9	
Q 9H8H3	Q9H8H3	TMT1A_HUMAN	TMT1A							2957.39	4628.61	4630.01
Q 9H8N7	Q9H8N7	ZN395_HUMAN	ZNF395	1580.51			1320.25				1633.97	998.416
Q 9H8V3	Q9H8V3	ECT2_HUMAN	ECT2					1115.29		1177.51		706.023
Q 9H8Y8	Q9H8Y8	GORS2_HUMAN	GORASP2	2441.19	1446.31	2071.17	1946.57	1896.71	2085.84	4190.08	1189.14	2827.86
Q 9H900	Q9H900	ZWILC_HUMAN	ZWILCH						1366.06	2199.67	3832.2	
Q 9H910	Q9H910	JUPI2_HUMAN	JPT2	3298.14		2302.41	4267.35	3289.64	5408.38	3227.62	3826.4	4647.42
Q 9H967	Q9H967	WDR76_HUMAN	WDR76	6337.43	10786.1		2094.71				3013.07	
Q 9H974	Q9H974	QTRT2_HUMAN	QTRT2	4181.28	4974.17	5011.07	2117.64	3963.87	2245.29	3281.11	2136.23	3125.77
Q 9H981	Q9H981	ARP8_HUMAN	ACTR8				760.651	1085.52	1171.26	860.358	521.624	880.558
Q 9H9A5	Q9H9A5	CNO10_HUMAN	CNOT10					1518.83	800.984	1396.23		908.8
Q 9H9A6	Q9H9A6	LRC40_HUMAN	LRRC40	7876.04	7434.59	6946.79	4667.74	5438.84	2708.87	5028.32	3399.37	5829.04
Q 9H9A7	Q9H9A7	RMI1_HUMAN	RMI1	10283.8	28595.9		4659.89	8034.69		1369	3130.86	
Q 9H9E3	Q9H9E3	COG4_HUMAN	COG4			7303.67	1944.68	2143.84	1684.84	2470.19	1566.22	3422.59
Q 9H9G7	Q9H9G7	AGO3_HUMAN	AGO3					11474.7	4755.7	4587.58		4774.6
Q 9H9H4	Q9H9H4	VP37B_HUMAN	VPS37B			1584.37	2084.46	3043.5	2989.72	2952.68	843.413	2084.8
Q 9H9P8	Q9H9P8	L2HDH_HUMAN	L2HGDH			1880.58	1809.3	1659.98	1132.53	7698.32	6035.68	8513.52
Q 9H9T3	Q9H9T3	ELP3_HUMAN	ELP3		3064.31	3974.77	8517.09	10718.3	12734.1	11419.7	8504.69	11950.7
Q 9H9Z2	Q9H9Z2	LN28A_HUMAN	LIN28A				8896.11	14927.1	13202.6			
Q 9HA77	Q9HA77	SYCM_HUMAN	CARS2							3900.05	2016.39	4026.52
Q 9HAC8	Q8WUN7;Q9HAC8	UBTD1_HUMAN	UBTD1				3204.65	8706.03	7097.63	11098.8	2148.1	9230.13
Q 9HAS0	Q9HAS0	NJMU_HUMAN	C17orf75				1043.31		1246.64	1312.44		1306.91
Q 9HAU0	Q9HAU0	PKHA5_HUMAN	PLEKHA5	11246.4	11017.6	10904.8	9214.18	8809.27	7767.68	2938.05	3326.33	3551.72
Q 9HAV4	Q9HAV4	XPO5_HUMAN	XPO5				5774.45	4180.2	4345.16	7410.36	6985.26	9801
Q 9HAV7	Q9HAV7	GRPE1_HUMAN	GRPEL1						3670.73	3347.31		2539.03
Q 9HB07	Q9HB07	MYG1_HUMAN	MYG1				2990.9	2114.34	7179.42	4223.43		2734.79
Q 9HB71	Q9HB71	CYBP_HUMAN	CACYBP	2229.3		3479.43	2346.85	2758.32	3561.74	3723.5	1754.53	3665.28
Q 9HB90;Q 9NQL2	Q9NQL2;Q9HB90	RRAGC_HUMAN;RRAGD_HUMAN	RRAGC;RRAGD				2239.66		2252.93	4585.02		4999.65
Q 9HBH1	Q9HBH1	DEFM_HUMAN	PDF							2299.56	1955.18	2812.22
Q 9HC07	Q9HC07	TM165_HUMAN	TMEM165			2682.68	2252.48		876.833	2790.09	2210.47	2654.19
Q 9HC35	Q9HC35	EMAL4_HUMAN	EML4	2179.85		2382.85	4879.08	3744.05	5398.95	3904.9	2936.25	3197.4
Q 9HC36	Q9HC36	MRM3_HUMAN	MRM3					850.29	474.341	1527.28	930.156	1320.92
Q 9HC38	Q9HC38	GLOD4_HUMAN	GLOD4			18154.9	21702.5		2402.52	6303.85		3724.35
Q 9HC84	Q9HC84	MUC5B_HUMAN	MUC5B			3012.7					3511.91	
Q 9HCC0	Q9HCC0	MCCB_HUMAN	MCCC2	967844	886827	1.07395E+06	496079	159907	156855	413533	758240	446840
Q 9HCE1	Q9HCE1	MOV10_HUMAN	MOV10	2243.67	3181.53	2154.96		4238.92	1192.88	1739.7	1342.64	2325.01
Q 9HCE6	Q9HCE6	ARGAL_HUMAN	ARHGEF10L					1970.74	2062.77			3005.57
Q 9HCE7	Q9HCE7	SMUF1_HUMAN	SMURF1				7822.25				5619.52	3569.13
Q 9HCH0	Q9HCH0	NCK5L_HUMAN	NCKAP5L						541.46			
Q 9HCJ3	Q9HCJ3	RAVR2_HUMAN	RAVER2							980.835	603.121	633.896
Q 9HCK8	Q9HCK8	CHD8_HUMAN	CHD8				1342.2			1004.42		784.016
Q 9HCP0	Q9HCP0	KC1G1_HUMAN	CSNK1G1					9743.48	4064.79	8192.31	2375.16	5708.81
Q 9HCU5	Q9HCU5	PREB_HUMAN	PREB	2355.81		2008.34	1859.36	2479.14	2172.44	3827.17	3742.32	3455.77
Q 9HCY8	Q9HCY8	S10AE_HUMAN	S100A14		6065.48	7362.27	3947.12	2938.29	3716.41	6909.15	4711.12	4145.04

Q9HD20	Q9HD20	AT131_HUMAN	ATP13A1	2054.73	1435.11	1700.62	1757	6682.07	3176.89	2333.38	1183.11	2479.61
Q9HD45	Q9HD45	TM9S3_HUMAN	TM9SF3	6563.56	6138.38	6536.66	5705.13	7856.39	8540.88	9297.41	7458.73	8661.61
Q9HD67	Q9HD67	MYO10_HUMAN	MYO10					7821.82	8217.7	6696.97	12637.1	11306.7
Q9HDC9	Q9HDC9	APMAP_HUMAN	APMAP	5750.58	3396.29	4170.23	3592.3	6118.91	6930.99	10442.5	9871.09	8617.44
Q9NNW5	Q9NNW5	WDR6_HUMAN	WDR6				2602.63	7417.2	5514.84	1163.46		5770
Q9NP55	Q9NP55	BPIA1_HUMAN	BPIFA1	7233.32	5720.18	12079.5	3541.79	3383.2	3616.42	4369.25	9875.51	2713.2
Q9NP61	Q9NP61	ARFG3_HUMAN	ARFGAP3					2379.1	2598.66	490.008		
Q9NP66	Q9NP66	HM20A_HUMAN	HMG20A	3387.89				1781.42	1591.08			
Q9NP72	Q9NP72	RAB18_HUMAN	RAB18									
Q9NP79	Q9NP79	VTA1_HUMAN	VTA1	3137.13		2943.85	4776.6	2783.66	3251.11	2186.3		2960.26
Q9NP81	Q9NP81	SYSM_HUMAN	SARS2	2954.06		3530.53	2888.63	3599.23	3164.05	2476.8	1930.31	3232.17
Q9NP97	Q9NP97;Q8TF09	DLRB1_HUMAN	DYNLRB1	2326.85		954.049	1390.23	3209.65	3408.08	1714.56	1538.73	2288.98
Q9NPA0	Q9NPA0	EMC7_HUMAN	EMC7		1423.69		1061.97		2712.7	2683.29		3718.03
Q9NPA3	Q9NPA3	MIIP1_HUMAN	MID1IP1			3843.58	3310.54	1800.76	2343.62	2333.11	3440.11	2272.01
Q9NPA8	Q9NPA8	ENY2_HUMAN	ENY2					5932.53	1789	1285.66		
Q9NPD3	Q9NPD3	EXOS4_HUMAN	EXOSC4	2778.33	3123.46	3188.16	1918.63	2373.85	5318.6	3678.43	1252.57	3936.82
Q9NPH2	Q9NPH2	INO1_HUMAN	ISYNA1					2665.36	5797.43			
Q9NPI6	Q9NPI6	DCP1A_HUMAN	DCP1A				648.699	2950.82	2586.94	723.203	594.466	557.03
Q9NPJ3	Q9NPJ3	ACO13_HUMAN	ACOT13					1230	2459.84	604.58	807.832	862.322
Q9NPQ8	Q9NPQ8;Q9NVN3	RIC8A_HUMAN	RIC8A				5831.46	5683.09	7737.25	6174.99	7813.85	6172.8
Q9NQ84	Q9NQ84	GPC5C_HUMAN	GPRC5C				4474.02	8528.82	7145.59	4543.31		4228.03
Q9NQA3	C4AMC7;Q6VEQ5;A8K0Z3;A8MWX3;Q9NQA3	WASH6_HUMAN	WASH6P	2632.02	2811.43	1755.29	2821.75	1793.08	3878.46	2652.38	1958.44	2845.92
Q9NQB0	Q9NQB0	TF7L2_HUMAN	TCF7L2				1968.41					
Q9NQC3	Q9NQC3	RTN4_HUMAN	RTN4	35297.6	18158.4	38939.2	24543.6	21754.5	25799	19323.8	17848.9	18478.9
Q9NQG5	Q9NQG5	RPR1B_HUMAN	RPRD1B				3587.08	3661.06	7355.79	1192.24		2209.86
Q9NQP4	Q9NQP4	PFD4_HUMAN	PFDN4	4277.08		2942.85	2428.52		2774.51	1962.84	1982.07	1833.62
Q9NQS1	Q9NQS1	AVEN_HUMAN	AVEN		623.337			1140.23	373.131	324.622		
Q9NQT4	Q9NQT4	EXOS5_HUMAN	EXOSC5					1111.28	1954.57	2458.79		1389.37
Q9NQT5	Q9NQT5	EXOS3_HUMAN	EXOSC3					696.615	824.249	704.344		561.394
Q9NQW6	Q9NQW6	ANLN_HUMAN	ANLN				668.746	671.263	704.738	543.023	304.564	289.142
Q9NQW7	Q9NQW7	XPP1_HUMAN	XPNPEP1	4326.16		3321.15	3826.43	5497.49	8941.76	4004.72	2258.27	4185.49
Q9NQX3	Q9NQX3	GEPH_HUMAN	GPHN					13100.7	5235.21	3180.04		
Q9NQX4	Q9NQX4	MYO5C_HUMAN	MYO5C				5348.14	11328.7	7000.99	25137.9	12896.1	24294.6
Q9NQZ2	Q9NQZ2	SAS10_HUMAN	UTP3							9204.52	17489.2	
Q9NR09	Q9NR09	BIRC6_HUMAN	BIRC6				1782.35		674.17	2672.54		2117.74
Q9NR12	Q9NR12	PDLI7_HUMAN	PDLIM7	17233.1	14910.1	15253.1	26129.5	8773.14	10023.8	6019.53	8649.08	6793.04
Q9NR30	Q9NR30	DDX21_HUMAN	DDX21	2724.18	3236.84	5552.5	7389.74	12225.9	9934.51	17963.8	18620.1	19278
Q9NR31	Q9NR31	SAR1A_HUMAN	SAR1A	11802.2	5801.25	16985.7	19800.1		2495.11	9564.64	14673	10958.4
Q9NR45	Q9NR45	SIAS_HUMAN	NANS		3099.78	5476.27	3817.74	3471.79	6368.3	3546.38		2847.16
Q9NR50	Q9NR50	EI2BG_HUMAN	EIF2B3	8759.82	8795.65	7641.55	11489.2	6559.43	7647.53	10457.8	10559.5	11572.1
Q9NR56	Q9NUK0;Q5VZF2;Q9NR56	MBNL1_HUMAN	MBNL1			1651.09	2125.62	3617.81	2710.42	1679.68	1681.23	2473.51
Q9NRA8	Q9NRA8	4ET_HUMAN	EIF4ENIF1				617.647	2836.63	1136.38	434.436		
Q9NRF8	Q9NRF8	PYRG2_HUMAN	CTPS2			2494.43	2835.01	2141.41	4305.48	4413.73	5389.72	5513.24
Q9NRG0	Q9NRG0	CHRC1_HUMAN	CHRAC1				934.634	584.925	604.474	444.749	1031.83	
Q9NRG9	Q9NRG9	AAAS_HUMAN	AAAS		4345.01	2173.58	8261.62	9581.16	2813.15	5530.37	6293.55	7788.89
Q9NRN7	Q9NRN7	ADPPT_HUMAN	AASDHPPT				4609.1	2513.22	8398.91	2622.07		3357.6
Q9NRP0	Q9NRP0	OSTC_HUMAN	OSTC	4035.96	3492.28	4407.1	2623.24	3970.35	3367.74	5753.77	2977.3	3881.96
Q9NRR4	Q9NRR4	RNC_HUMAN	DROSHA		2592.6			1473.95		815.675		376.718
Q9NRR6	Q9NRR6	INP5E_HUMAN	INPP5E									
Q9NRR8	Q9NRR8	C42S1_HUMAN	CDC42SE1					1654.48	1002.99	2680.16		2638.43
Q9NRW3	Q9NRW3	ABC3C_HUMAN	APOBEC3C	3405.59	7504.44	2279.96	1948.09	3159.06		2138.91	1756.2	1208.58
Q9NRW7	Q9NRW7	VPS45_HUMAN	VPS45				1866.17	3520.47	2047.85	1318.07		1800.14
Q9NRX1	Q9NRX1	PNO1_HUMAN	PNO1	3057.84	3382.05		3329.24	1459.6	1020.5	1952.04	3407.33	1990.78
Q9NRX2	Q9NRX2	RM17_HUMAN	MRPL17			2906.58	2207.34			4208.77	3548.99	5245.51
Q9NRY2	Q9NRY2	SOSSC_HUMAN	INIP						919.385	2652.56	3090.62	1983.9
Q9NRY5	Q9NRY5	F1142_HUMAN	FAM114A2					2515.53	1828.52	306.88		266.415
Q9NRZ9	Q9NRZ9	HELLS_HUMAN	HELLS				1878.61	2535.19	1851.63	5163.86	6242.56	4912.71
Q9NS93	Q9NS93	TM7S3_HUMAN	TM7SF3		3567.68					3739.97	814.532	2533.69
Q9NSB2	Q9NSB2	KRT84_HUMAN	KRT84	4519.55	2029.85		5146.86				4685.2	2638.13
Q9NSB4	Q9NSB4	KRT82_HUMAN	KRT82			4308.04					4817.61	
Q9NSD9	Q9NSD9	SYFB_HUMAN	FARSB	4177.8	2558.84	3496	4764.29	8682.12	6006.3	4059.75	3932.23	6106.26
Q9NSE4	Q9NSE4	SYIM_HUMAN	IARS2				4958.81	2885.97	3302.69	3653.5	3189.5	4011.45
Q9NSI2	Q9NSI2	SLX9_HUMAN	SLX9	9445.95	4771.94	5645.92	5234.85	14207.2	5886.17	7990.06	5658.99	8825.91
Q9NSU2	Q9NSU2	TREX1_HUMAN	TREX1	20222.1	46103.5		2469.22	1595.65		12548.1		
Q9NSV4	Q9NSV4	DIAP3_HUMAN	DIAPH3	10899.4	4996.74	5547.17	7061.86	6957.48		4166.79	10320.1	4560.41
Q9NT62	Q9NT62	ATG3_HUMAN	ATG3	2925.45		2044.59	2634.2		13892.9	6836.34	1916.97	3206.76
Q9NTI5	Q9NTI5	PDS5B_HUMAN	PDS5B	779.482			717.768	1843.61	827.088	805.938	763.623	596.891
Q9NTJ3	Q9NTJ3	SMC4_HUMAN	SMC4	3844.34	3546.29	2598.58	6299.26	6684.52	5751.03	7928.55	7446.86	8071.71
Q9NTJ5	Q9NTJ5	SAC1_HUMAN	SACM1L	7143.29	8531.14	7655.32	5143.19	4021.61	4899.65	6465.01	5664.56	4753.72
Q9NTK5	Q9NTK5	OLA1_HUMAN	OLA1	6313.4	2747.46	9929.8	10951.8	5701.74	7090.49	14438.5	7114.47	9820.58
Q9NTX5	Q9NTX5	ECHD1_HUMAN	ECHDC1		510.795			1201.05	558.669	585.762		1914.93
Q9NUD5	Q9NUD5	ZCHC3_HUMAN	ZCCHC3				477.824	11032.5	1861.77	1115.61	1355.94	
Q9NUL3	Q9NUL3	STAU2_HUMAN	STAU2	3481.95	3612.8	2899.96	1959.56	5752.94	2440.76	3345.59	2559.89	2761.74
Q9NUL7	Q9NUL7	DDX28_HUMAN	DDX28	2765.93	2934.19	2954.45	3814.16	934.605	617.405	2589.59	3387.85	3715.11
Q9NUM4	Q9NUM4	T106B_HUMAN	TMEM106B	3644.7	2301.6	3766.96	1915.83	2814.71	4090.94	2320.72		3593.51
Q9NUP9	Q9NUP9	LIN7C_HUMAN	LIN7C					740.73	990.711	1817.67	647.398	1612.82
Q9NUQ3	Q9NUQ3	TXLNG_HUMAN	TXLNG					521.246	336.808		1328.11	533.043
Q9NUQ6	Q9NUQ6	SPS2L_HUMAN	SPATS2L					3671.97	2040.12	2491.45		3059.62
Q9NUQ8	Q9NUQ8	ABCF3_HUMAN	ABCF3	6250.64	6640.6	4016.01	6069.81	3472.17	2368.5	4309.81	6852.97	4498.9
Q9NUQ9	Q9NUQ9	CYRIB_HUMAN	CYRIB				2489.6				862.586	573.707
Q9NUU7	Q9NUU7	DD19A_HUMAN	DDX19A	1708.43		2210.92	5825.33	1486.74	2095.47	3689.99	6795.79	3329.82
Q9NV70	Q9NV70	EXOC1_HUMAN	EXOC1				3890.55	6237.82	5350.53	3844.04	3610.22	4092.28
Q9NVA2	Q9NVA2;Q14141;Q92599	SEP11_HUMAN	SEPTIN11	6492.65		5360.97	9692.28	4415.55	9012.97	3171.21	3166.85	4762.38
Q9NVC6	Q9NVC6	MED17_HUMAN	MED17				864.989	671.238	579.862	1532.24	908.745	947.538
Q9NVE7	Q9NVE7	PANK4_HUMAN	PANK4				1905.39			766.374	1121.33	
Q9NVH1	Q9NVH1	DJC11_HUMAN	DNAJC11	2241.76	2240.38				911.386	1100.66		1363.06
Q9NVH6	Q9NVH6	TMLH_HUMAN	TMLHE									
Q9NVI1	Q9NVI1	FANCI_HUMAN	FANCI				3447.2	2043.8	1707.4	2809.07	4995.36	3097.67
Q9NVI7	Q9NVI7;Q5T9A4	ATD3A_HUMAN	ATAD3A	9122.32	12312.4	9184.12	1576.85	6179.72	2086.85	8758.14	5232.15	6870.19
Q9NVJ2	Q9NVJ2;Q96BM9	ARL8B_HUMAN	ARL8B	6474.69	4406.35	10146.1		3074.52	3446.29	3939.35	3453.19	4242.14
Q9NVM9	Q9NVM9	INT13_HUMAN	INTS13	9121.32	10100.5	6470.77	7996.73	7493.97	5205.62	17447.2	17975.5	16496.5
Q9NVN8	Q9NVN8	GNL3L_HUMAN	GNL3L	4251.68	6286.96	2528.4	4054.99	2855.39	2700.99	5742.6	6479.92	6628.29
Q9NVT9	Q9NVT9	ARMC1_HUMAN	ARMC1					2192.39	1638.21	674.138		
Q9NVU0	Q9NVU0	RPC5_HUMAN	POLR3E	7165	9072.47		2325.02		981.952			1543.6
Q9NVU7	Q9NVU7	SDA1_HUMAN	SDAD1		1526.7			896.207	1111.64	1613.73		1235.47
Q9NVV4	Q9NVV4	PAPD1_HUMAN	MTPAP	3484.8		3003.11	1671.11	2212.39	1496.86	4171.49	4148.71	5358.3
Q9NW08	Q9NW08	RPC2_HUMAN	POLR3B	6407.45	7089.29		2441.87	2989.64	2747.1	2361.11	1083.61	2351.06
Q9NWB1	Q9NWB1	DMAC2_HUMAN	DMAC2				1633.24	3880.64	2199.21	2535.86	2397.86	2414.47
Q9NWQ8	Q9NWQ8	PHAG1_HUMAN	PAG1				555.791	1099.06	1051.15	600.016		638.539
Q9NWV4	Q9NWV4	CZIB_HUMAN	CZIB						1607.01			
Q9NWZ5	Q9NWZ5	UCKL1_HUMAN	UCKL1				1677.33	4379.31	3182.66	2088.22	2540.94	2534.68
Q9NX05	Q9NX05	F120C_HUMAN	FAM120C		3371.54	2095.2		3575.32	552.475	4527.6	2068.18	2068.95
Q9NX08	Q9NX08	COMD8_HUMAN	COMMD8			2278.54	1153.98		1711.26	1278.13	843.565	1680.78
Q9NX20	Q9NX20	RM16_HUMAN	MRPL16	3641.42	5080.14	3754.92	1916.5	6390.72	4150.04	7296.5	4500.84	6564.36
Q9NX57	Q9NX57	RAB20_HUMAN	RAB20							5484.16	1387.7	4918.6
Q9NX62	Q9NX62	IMPA3_HUMAN	BPNT2					1689.85	1869.8	1417.52		1021.23
Q9NX63	Q9NX63	MIC19_HUMAN	CHCHD3	17040.6	23679.9	18246.3		6680.93	5743.92	5564.79		5667.77
Q9NXC5	Q9NXC5	MIO_HUMAN	MIOS					1918.83	1619.21	928.914		834.225
Q9NXE4	Q9NXE4	NSMA3_HUMAN	SMPD4					702.506	1136.73	1218.6		1091.24
Q9NXG2	Q9NXG2	THUM1_HUMAN	THUMPD1						2521.13	588.285		
Q9NXH8	Q9NXH8	TOR4A_HUMAN	TOR4A	4712.24	5742.23	3989.04	4892.58	9438.08	6297.62	3738.02	2653.37	3722.55
Q9NXH9	Q9NXH9	TRM1_HUMAN	TRMT1				1196		3097.21	2546.92		1922.18
Q9NXK8	Q9NXK8	FXL12_HUMAN	FBXL12				898.625	6540.47	2216.62			
Q9NXR7	Q9NXR7	BABA2_HUMAN	BABAM2		1273.93	1302.75		6913.4	4221.45	5205.42		3964.6
Q9NXV6	Q9NXV6	CARF_HUMAN	CDKN2AIP		8717.21			1483.61		2228.13		
Q9NXW2	Q9NXW2	DJB12_HUMAN	DNAJB12	1504.45	2362.99	1083.04	894.313	2548.44	1556.3	6073.05	2382.93	6229.7
Q9NXZ2	Q9NXZ2	DDX43_HUMAN	DDX43				1946.21	2091.19	1599.16			
Q9NY12	Q9NY12	GAR1_HUMAN	GAR1					4508.67				
Q9NY33	Q9NY33	DPP3_HUMAN	DPP3				1727.28	537.418	989.976	561.577	1005.85	408.338
Q9NY35	Q9NY35	CLDN1_HUMAN	CLDND1				3325.98	3726.19	8417.78	8317.9	2458.09	9772.81
Q9NY65	Q9NY65	TBA8_HUMAN	TUBA8	55078.2	23703.5	38386.1	54456.8	12708.9	18258.2	28308.3	80419.4	63337.1
Q9NY93	Q9NY93	DDX56_HUMAN	DDX56		4584.9		1109.17	7740.93	4233.3	2847.07	1651.02	1478.03
Q9NYB9	Q9NYB9	ABI2_HUMAN	ABI2				2286.24	3023.78	3125.23	3643.32		4116.87
Q9NYF8	Q9NYF8	BCLF1_HUMAN	BCLAF1	3787.85	7311.76	1998.73	735.225	3850.3	883.995	1578.02	957.632	1445.48
Q9NYJ1	Q9NYJ1	COA4_HUMAN	COA4					1210.22	3105.52	1005.59		1811.11
Q9NYL2	Q9NYL2	M3K20_HUMAN	MAP3K20	8077.9	10736.1	6334.2	7774.77	5090.14	5298.42	5830.96		6528.6
Q9NYL4	Q9NYL4	FKB11_HUMAN	FKBP11			3275.05				6367.28	7683.53	7762.61
Q9NYL9	Q9NYL9	TMOD3_HUMAN	TMOD3	7516.71	8496.76	6782.12	32807.9	82924.6	44662.3	56526.7	20320.9	44172.3
Q9NYU2	Q9NYU2	UGGG1_HUMAN	UGGT1	2965.81	2503.61	1743.65	1572.28	2380.93	4275.34	2258.35		623.793
Q9NYV6	Q9NYV6	RRN3_HUMAN	RRN3					383.496		1247.46	981.186	1258.21
Q9NYY8	Q9NYY8	FAKD2_HUMAN	FASTKD2	1137.77	2634.44	3811.78	4492.72	2232.51	3094.78	5331.93	7788.62	4902.78
Q9NYZ3	Q9NYZ3	GTSE1_HUMAN	GTSE1				4706.68	3557.3		4410.4	2866.03	4374.99
Q9NZ01	Q9NZ01	TECR_HUMAN	TECR	5545.05	5799.47	7699.73	3052.82	3745.22	3917.47	8263.93	4904.69	10969.4
Q9NZ08	Q9NZ08	ERAP1_HUMAN	ERAP1	2553.4	3664.65	2592.81	1788.81	7371.66	4514.81	2980.97	1965.26	3599.05
Q9NZ09	Q9NZ09	UBAP1_HUMAN	UBAP1	3448.3	3925.77	3483.62		1547.24	1591.5			1870.77
Q9NZ63	Q9NZ63	TLS1_HUMAN	C9orf78					2227.61	2807.84			
Q9NZB2	Q9NZB2	F120A_HUMAN	FAM120A	11198.7	6476.61	4651.55	7832.81	12241.9	4402.38	4150.22	5973.95	4574.88
Q9NZC4	Q9NZC4	EHF_HUMAN	EHF							779.256		724.731
Q9NZC9	Q9NZC9	SMAL1_HUMAN	SMARCAL1	12610.8	33344.6	3086.55	7699.38	10832	2409.76	783.189	1349.66	
Q9NZH8	Q9NZH8	IL36G_HUMAN	IL36G				4177.39				4168.11	
Q9NZI7	Q9NZI7	UBP1_HUMAN	UBP1					1731.48		340.632	701.729	
Q9NZI8	Q9NZI8	IF2B1_HUMAN	IGF2BP1	24140.7	8069.17	13492.3	33501.8	74690.2	28813.6	9231.98	13685.9	9969.07
Q9NZL4	Q9NZL4	HPBP1_HUMAN	HSPBP1				2850.87	3624.47	6074.21	4243.29	2095.72	2770.11
Q9NZL9	Q9NZL9	MAT2B_HUMAN	MAT2B				2743.91		686.972		2845.34	
Q9NZM1	Q9NZM1	MYOF_HUMAN	MYOF	11296.4	7348.29	9934.97	13567.8	17192.2	14823.5	22375.3	12238.2	18811.1
Q9NZN4	Q9NZN4	EHD2_HUMAN	EHD2	17341.5	18346.4	17417.3	9663.67	7813.93	6532.49	4701.64	4470	8491.25
Q9NZN5	Q9NZN5	ARHGC_HUMAN	ARHGEF12				2343.09		2189.21	3313.82	4570.42	2798.68
Q9NZN8	Q9NZN8	CNOT2_HUMAN	CNOT2	1199.25			3378.28	804.73	1100.18	2332.51	3444.2	2121.78
Q9NZR1	Q9NZR1	TMOD2_HUMAN	TMOD2							1412.12		2491.91
Q9NZT1	Q9NZT1	CALL5_HUMAN	CALML5	8963.61	8015.8	16222.8	5615.01	6782.88	4246.61	3535.8	8773.15	6577.37
Q9NZT2	Q9NZT2	OGFR_HUMAN	OGFR				1778.41	1310.75	2173.29	1808.14		2627.66
Q9NZW5	Q9NZW5	PALS2_HUMAN	PALS2				1446.7		1118.89	2804.01	2892.93	3914.5
Q9NZZ3	Q9NZZ3	CHMP5_HUMAN	CHMP5	4335.28	1522.6	2820.42	3267.5	2372.28	3999.17	1610.48	2003.77	1364.7
Q9P015	Q9P015	RM15_HUMAN	MRPL15	3148.98		2596.34	2196.18	1933.2	1451.06	4025.97	2843.46	3846.88
Q9P016	Q9P016	THYN1_HUMAN	THYN1		13551.2			1044.55	1745.63			
Q9P035	Q9P035	HACD3_HUMAN	HACD3				1925.61	3081.04	3464.42	2083.46	1555.84	3084.27
Q9P0I2	Q9P0I2	EMC3_HUMAN	EMC3	4616.91	4842.12	4121.98	2504.87		2883.45	8341.28	2398.62	7209
Q9P0J0	Q9P0J0	NDUAD_HUMAN	NDUFA13					2187.81	1229.59	2773.23	1100.06	4365.86
Q9P0J1	Q9P0J1	PDP1_HUMAN	PDP1		1017.28	14337.4	1476.61	2086.54	3533.56	1510.3		
Q9P0K7	Q9P0K7	RAI14_HUMAN	RAI14	43949.6	53046	35806.8	84187.5	156808	109809	9364.32	8515.17	9675.2
Q9P0L0	Q9P0L0	VAPA_HUMAN	VAPA	5067.65	4538.95	6240.05	4829.25	5693.14	4619.84	9287.4	4251.63	8026.56
Q9P0V9	Q9P0V9	SEP10_HUMAN	SEPTIN10				2259.5	713.662	1217.55	570.606		
Q9P1Y5	Q9P1Y5	CAMP3_HUMAN	CAMSAP3							4101.02	1488.36	2295.85
Q9P1Z2	Q9P1Z2	CACO1_HUMAN	CALCOCO1	3780.9	2593.08	4858.29	1558.17	1550.05	1578.23	1696.63	849.83	2200.56
Q9P206	Q9P206	K1522_HUMAN	KIAA1522	12950.4	13019	8578.24	9584.38	30800.4	16454.4	14220.5	7886.14	15329.6
Q9P258	Q9P258	RCC2_HUMAN	RCC2	3372.69	3960.88	3480.13	8622.68	5274.51	6290.14	6898.43	6680.46	8022.05
Q9P260	Q9P260	RELCH_HUMAN	RELCH			754.394	2617.93	647.949	877.892	542.228	1039.44	585.447
Q9P265	Q9P265	DIP2B_HUMAN	DIP2B	4433.14	3228.9		3910.85			1175.6		
Q9P270	Q9P270	SLAI2_HUMAN	SLAIN2	1067.51				1002.85	730.128			
Q9P287	Q9P287	BCCIP_HUMAN	BCCIP					2704.63	2140.33	1977.25	339.265	2341.03
Q9P289	Q9Y6E0;Q9P289	STK26_HUMAN	STK26		2646.09	1627.18	3068.97	1585.82	2110.18	2801.32	2868.24	3280.77
Q9P2D3	Q9P2D3	HTR5B_HUMAN	HEATR5B				1060.04		281.946	551.108	1090.95	
Q9P2E9	Q9P2E9	RRBP1_HUMAN	RRBP1	7660.51	4651.82	4815.31	3919.58	4305.03	3000.8	5049.54	3005.31	4544.52
Q9P2I0	Q9P2I0	CPSF2_HUMAN	CPSF2		1291.18			1225.76	916.773	1137.08		1122.24
Q9P2J5	Q9P2J5	SYLC_HUMAN	LARS1	4896.58	3870.33	3888.65	11768.9	8242.07	7651.98	6854.18	4316.33	6416.84
Q9P2K3	Q9P2K3	RCOR3_HUMAN	RCOR3									
Q9P2K5	Q9P2K5	MYEF2_HUMAN	MYEF2									
Q9P2K8	Q9P2K8	E2AK4_HUMAN	EIF2AK4	1054.03			1958.6	2087.69	2206.63	777.25		872.396
Q9P2M7	Q9P2M7	CING_HUMAN	CGN				2985.91	4324.23	2608.04	1800.36		1447.36
Q9P2N5	Q9P2N5	RBM27_HUMAN	RBM27	5121.83	4542.57	5763.47	3717.22	6459.74	1544.44	2558.82	2152.01	2192.33
Q9P2R3	Q9P2R3	ANFY1_HUMAN	ANKFY1	6211.54	9909.51	5310.02	3481.63		4154.86	4485.26	2579.78	
Q9P2R7	Q9P2R7	SUCB1_HUMAN	SUCLA2	4528.97	3498.89	3131.44	2703.5	6919.71	5384.71	1744.98	5244.13	2234.09
Q9P2T1	Q9P2T1	GMPR2_HUMAN	GMPR2				2926.91	1152.49	1353.38	590.303		
Q9P2Y4	Q9P2Y4	ZN219_HUMAN	ZNF219									
Q9UBB4	Q9UBB4	ATX10_HUMAN	ATXN10	3144.43	3394.89	3741.63	5199.97	5973.16	2782.64	3362.11	5342.25	3183.98
Q9UBB5	Q9UBB5	MBD2_HUMAN	MBD2	3355.87			1397.44	791.687	919.191	1298.04		1108.78
Q9UBB6	Q9UBB6	NCDN_HUMAN	NCDN		2352.02		3350.93	3041.7	2962.01	3782.21	2923.58	3409.74
Q9UBB9	Q9UBB9	TFP11_HUMAN	TFIP11		3946.72		3876.38	2813.13	1440.45	2746.57	4739.21	2213.02
Q9UBC2	Q9UBC2	EP15R_HUMAN	EPS15L1				1829.17	796.864	1055.45	479.11		558.25
Q9UBC9	Q9UBC9	SPRR3_HUMAN	SPRR3								2929.31	
Q9UBD5	Q9UBD5	ORC3_HUMAN	ORC3				1060.72	2354.44	1045.18	1489.36	743.337	897.362
Q9UBE0	Q9UBE0	SAE1_HUMAN	SAE1			6019.24		2409.02	7798.3	5250.35		5406.12
Q9UBF2	Q9UBF2	COPG2_HUMAN	COPG2	4572.45	3073.04	5324.76	10498.5	5873.91	3835.24	6282	7771.67	6881.27
Q9UBG3	Q9UBG3	CRNN_HUMAN	CRNN				1347.62				2143.21	
Q9UBI1	Q9UBI1	COMD3_HUMAN	COMMD3				239.846	1102.49	1194.67	1082.72		1266.33
Q9UBI6	Q9UBI6	GBG12_HUMAN	GNG12	5951.79	8820.04	7338.33	38388.6	75142.7	50743.7	26714.6	13685	36556.8
Q9UBM7	Q9UBM7	DHCR7_HUMAN	DHCR7							5235.59	3407.5	6692.33
Q9UBP6	Q9UBP6	TRMB_HUMAN	METTL1				619.619	446.529	2312.42	785.435	231.453	637.426
Q9UBQ0	Q9UBQ0	VPS29_HUMAN	VPS29						1742.95			787.239
Q9UBQ5	Q9UBQ5	EIF3K_HUMAN	EIF3K			2037.69	1899.26	1954.22	4691.37	1389.96	2358.37	1307.11
Q9UBQ7	Q9UBQ7	GRHPR_HUMAN	GRHPR					2776.67	4531.19	2898.93		3169.63
Q9UBR2	Q9UBR2	CATZ_HUMAN	CTSZ									
Q9UBS4	Q9UBS4	DJB11_HUMAN	DNAJB11	1098.56	3138.78	3764.49	3792.07	4412.98	4811.43	5132.66	2793.36	4984.34
Q9UBT2	Q9UBT2	SAE2_HUMAN	UBA2			2552.52	1445.19	1781.45	5372.28	2534.06		2525.03
Q9UBU6	Q9UBU6	FA8A1_HUMAN	FAM8A1							737.24	1405.79	675.656
Q9UBV2	Q9UBV2	SE1L1_HUMAN	SEL1L			9001.77	1315.45	3908.99	1476.3	1831.25	1305.66	3178.99
Q9UBV8	Q9UBV8	PEF1_HUMAN	PEF1					4074.06	3514.65	1623.58		
Q9UBX3	Q9UBX3	DIC_HUMAN	SLC25A10							533.025		690.061
Q9UBX7	Q9UBX7	KLK11_HUMAN	KLK11	50521.5	60739.2	61858.6	87249.6	10141.8	20413	39402.1	28062.4	33926.9
Q9UDR5	Q9UDR5	AASS_HUMAN	AASS	906.721		1279.38	1477.02	2271.24	2774.94	309.441		
Q9UDY2	Q9UDY2	ZO2_HUMAN	TJP2				8154.98	10943.8	9723.25	4169.43	6747.3	5956.1
Q9UET6	Q9UET6	TRM7_HUMAN	FTSJ1		1163.9		1327.11	310.215	669.176	1107.33		1666.84
Q9UEW8	Q9UEW8	STK39_HUMAN	STK39	16871.9	9617.59	18228.2	4826.35		5448.34	6438.89	9027.83	8449.47
Q9UFC0	Q9UFC0	LRWD1_HUMAN	LRWD1		1146.43		3568.15	2720.83	3788.36	1690.2	1608.21	1532.8
Q9UFW8	Q9UFW8	CGBP1_HUMAN	CGGBP1				2345.86			1631.55	2177.6	
Q9UG63	Q9UG63	ABCF2_HUMAN	ABCF2	13981.5	16222.5	12489.9	11164.3	14206.4	11784.8	17490.2	17929.4	16458.2
Q9UGI8	Q9UGI8	TES_HUMAN	TES				5123.37	7140.69	4385.82	5142.1	3218.5	2734.02
Q9UGJ1	Q9UGJ1	GCP4_HUMAN	TUBGCP4				3891.62	3214.82	1957.33	3696.29	3519.04	4073.37
Q9UGM3	Q9UGM3	DMBT1_HUMAN	DMBT1	9393.49		17010.6	1957.93			1558.65	3403.47	2195.81
Q9UGN5	Q9UGN5	PARP2_HUMAN	PARP2	16735.4	25735.9		9899.79	11146.3	3668.92	3832.04	9749.53	
Q9UGP4	Q9UGP4	LIMD1_HUMAN	LIMD1	3361.34	5260.67	4088.32	1340.34	1798.74	2904.38	2456.96	1176.02	1879.59
Q9UGP8	Q9UGP8	SEC63_HUMAN	SEC63		4947.1	3904.95	1820.99	2286.79	2686.51	4339.6	1536.18	4022.38
Q9UGR2	Q9UGR2	Z3H7B_HUMAN	ZC3H7B					2315.56	1687.01	874.733		958.406
Q9UH62	Q9UH62	ARMX3_HUMAN	ARMCX3				1119.97	1881.22		1546.37	646.237	2074.83
Q9UH99	Q9UH99	SUN2_HUMAN	SUN2	8388.25	10402.7	8970.15	5909.9	8176.1	6868.8	2403.82	2814.97	2091.68
Q9UHA4	Q9UHA4	LTOR3_HUMAN	LAMTOR3	2988.26			4283.24	4368.43	8119.4	22842.8	9228.26	6388.43
Q9UHB6	Q9UHB6	LIMA1_HUMAN	LIMA1	5783.32	7596.27	6833.55	12208.6	90884.8	58562.3	44287.5	7671.92	34814.7
Q9UHB9	Q9UHB9	SRP68_HUMAN	SRP68	4445.37	6125.65	3908.87	6663.97	6530.12	5277.64	4598.76	4768.07	4548.38
Q9UHD1	Q9UHD1	CHRD1_HUMAN	CHORDC1			2035.87	7289.75	8686.36	16174.6	6701.98	2192.11	6877.66
Q9UHD2	Q9UHD2	TBK1_HUMAN	TBK1		1342.71		840.166	792.464	825.81	1272.49		1093.05
Q9UHD8	Q9UHD8	SEPT9_HUMAN	SEPTIN9	13764.5	13356.1	15719.8	20265.6	12312.8	13255.8	12014.6	14385.5	12089.6
Q9UHI6	Q9UHI6	DDX20_HUMAN	DDX20		5120.08		5870.43	6161.84	4109.43	6062.97	5346.44	6943.84
Q9UHJ6	Q9UHJ6	SHPK_HUMAN	SHPK					2313.77	2135.67	1319.07	976.116	
Q9UHK6	Q9UHK6	AMACR_HUMAN	AMACR							2761.33	1678	1605.83
Q9UHN1	Q9UHN1	DPOG2_HUMAN	POLG2	8640.25	21652		12059.8	11003.6		4425.59		
Q9UHN6	Q9UHN6	CEIP2_HUMAN	CEMIP2	1197.84				1572.89	1224.06	4628.36	1628.42	3810.71
Q9UHR6	Q9UHR6	ZNHI2_HUMAN	ZNHIT2			3313.11	6053.5		2038.86	6678.58	10095.6	4853.8
Q9UHV9	Q9UHV9	PFD2_HUMAN	PFDN2			4744.48	5684.58		2990.74	3926.53	5761.45	4635.58
Q9UHX1	Q9UHX1	PUF60_HUMAN	PUF60	12725.2	17808.4	10865	7446.23	40251.7	16892	16877.2	5634.52	9241.81
Q9UHY7	Q9UHY7	ENOPH_HUMAN	ENOPH1						1373.45			
Q9UI10	Q9UI10	EI2BD_HUMAN	EIF2B4	9501.84	7167.75	6175.23	10071.2	10219.8	4183.55	9398.25	10436.9	7677.17
Q9UI12	Q9UI12	VATH_HUMAN	ATP6V1H	3608.54		2904.59	5935.9	5029.97	2680.57	3585.49	8778.12	4677.72
Q9UI26	Q9UI26	IPO11_HUMAN	IPO11				4885.78	2104.51	2033.32	2664.41	3773.05	3192.22
Q9UI30	Q9UI30	TR112_HUMAN	TRMT112	7718.85	3304.36		4936.22	5129.94	4699.54	6433.17	4718.72	5781.69
Q9UI42	Q9UI42	CBPA4_HUMAN	CPA4	4247.2	1611.46	8959.76	3766.29	4676.43	5708.25	4414.57	3811.81	3346.3
Q9UIA9	Q9UIA9	XPO7_HUMAN	XPO7	3684.15	2434.96	2157.16	5877.81	5434.71	6838.61	6106.48	5591.26	5194.53
Q9UID3	Q9UID3	VPS51_HUMAN	VPS51				2397.43	2514.44	2907.67	3961.07	2222.88	3896.18
Q9UII2	Q9UII2	ATIF1_HUMAN	ATP5IF1				1604.1				3892.01	
Q9UII4	Q9UII4	HERC5_HUMAN	HERC5					3329.34	694.345			
Q9UIJ7	Q9UIJ7	KAD3_HUMAN	AK3		3476.66			5674.85	2534.2	4836.78		3579.07
Q9UIQ6	Q9UIQ6	LCAP_HUMAN	LNPEP				1402.44	2920.13	3732.57	1225.74	1171.53	1592.65
Q9UIV1	Q9UIV1	CNOT7_HUMAN	CNOT7					2543.32	1855.82	964.348		2837.12
Q9UJ70	Q9UJ70	NAGK_HUMAN	NAGK	4745.87		4511.13	2526.92			1140.74	2514.09	877.366
Q9UJA3	Q9UJA3	MCM8_HUMAN	MCM8					1304.41		1711.85	1749.36	2217.07
Q9UJA5	Q9UJA5	TRM6_HUMAN	TRMT6				1494.87	6551.22	3865.49	4194.56	1555.25	3910.8
Q9UJC3	Q9UJC3	HOOK1_HUMAN	HOOK1							3711.29	1629.06	3523.2
Q9UJC5	Q9UJC5	SH3L2_HUMAN	SH3BGRL2					2674	495.806	7576.63	806.943	1311.43
Q9UJF2	Q9UJF2	NGAP_HUMAN	RASAL2				2902.14	982.169	799.903		974.344	538.516
Q9UJS0	Q9UJS0	S2513_HUMAN	SLC25A13	2737.25	2648.69	1999.62	3075.53	2530.63	2649.71	4798.05	5016.67	5657.96
Q9UJU6	Q9UJU6	DBNL_HUMAN	DBNL	3055.56			4188.84	4948.42	4303.82	2210.45		2132.57
Q9UJV9	Q9UJV9	DDX41_HUMAN	DDX41	5232.25	4579.77	5381.17	5354.15	4776.03	2463.16	6452.33	10527.2	6869.15
Q9UJX3	Q9UJX3	APC7_HUMAN	ANAPC7	1187.36	1692.36	1092.05	1117.04	3371.93	2385.67	2801.77	1986.81	1216.88
Q9UJX4	Q9UJX4	APC5_HUMAN	ANAPC5		1215.11		2112.82	4177.09	3312.85	4709.64	2299.33	4948.92
Q9UJX6	Q9UJX6	ANC2_HUMAN	ANAPC2		699.435	1117.81	1282.01	1290.03	1573.93	996.956	438.644	646.98
Q9UJZ1	Q9UJZ1	STML2_HUMAN	STOML2	3580.22	2427.79	2686.91	2219.49	1843.52	1157	3709.42	3524.2	3529.69
Q9UK76	Q9UK76	JUPI1_HUMAN	JPT1	1928.09		2293.32	2378.21	7584.03	15122.6	2533.9	3883.9	4084.43
Q9UKD2	Q9UKD2	MRT4_HUMAN	MRTO4	9542.56	13157.1	8544.93	8845.79	12694.4	8083.71	12505.7	14518.5	12358.9
Q9UKE5	Q9UKE5	TNIK_HUMAN	TNIK			18177.7		3295.27	2339.64	3316.66		4315.91
Q9UKF6	Q9UKF6	CPSF3_HUMAN	CPSF3				877.146	6049.49	3444.8	4083.36	3222.73	1756.55
Q9UKG1	Q9UKG1	DP13A_HUMAN	APPL1				1785	842.043	916.175	1062.49	2443.85	1534.61
Q9UKK3	Q9UKK3	PARP4_HUMAN	PARP4	8110.83	15079.5	3380.49	2386.19	5983.29	3788.84	1898.47		2028.58
Q9UKK9	Q9UKK9	NUDT5_HUMAN	NUDT5	4890.98	1920.8	4761.59	3208.89	2402.7	4839.01	5170.96	2987.26	4343.69
Q9UKL0	Q9UKL0	RCOR1_HUMAN	RCOR1	3317.81			4216.32	1497.15		2506.8	5698.62	1445.73
Q9UKM9	Q9UKM9	RALY_HUMAN	RALY	1422.04	774.341	974.647	1710.55	7122.44	1543.77	3039.76	2144.25	3656.6
Q9UKN1	Q9UKN1	MUC12_HUMAN	MUC12	137272	113225	99101.5	104028	22813.1	23242.3	47822.5	113067	39328.6
Q9UKN8	Q9UKN8	TFIC4_HUMAN	GTF3C4				2176.24	3083.93	2226.65	1608.46		2184.42
Q9UKS6	Q9UKS6	PACN3_HUMAN	PACSIN3			3480.71	10630.4	17874.2	15183.3	26746.2	7053.08	26058.9
Q9UKU7	Q9UKU7	ACAD8_HUMAN	ACAD8				976.399		884.539	1648.8	1322.23	1818.19
Q9UKU9	Q9UKU9	ANGL2_HUMAN	ANGPTL2	4718.95	1929.71	3426.27						
Q9UKV3	Q9UKV3	ACINU_HUMAN	ACIN1		3413.14			11795.8	1611.66	2300.52	1234.88	
Q9UKV8	Q9UKV8	AGO2_HUMAN	AGO2	5021.8			3806.84	4890.73	1488.69	475.093	4217.57	
Q9UKY7	Q9UKY7	CDV3_HUMAN	CDV3	12619.3	7331.04	16451.3	7839.42	6988.63	30636.4	13679.1	5365.05	16292.2
Q9UL03	Q9UL03	INT6_HUMAN	INTS6	4470.38	7316.15	4214.49	1906.8	3978.75	1907.72	4488.44	3495.01	4255.54
Q9UL15	Q9UL15	BAG5_HUMAN	BAG5						639.251	1237.84		1313.67
Q9UL46	Q9UL46	PSME2_HUMAN	PSME2		4761.31	4703.31	2878.57	2502.2	4531.67	3857.61	2988.38	4155.72
Q9UL54	Q9UL54	TAOK2_HUMAN	TAOK2		2617.36	5893.98	2498.71	2990.71	1652.37	1586.16	2728.59	2381.63
Q9ULA0	Q9ULA0	DNPEP_HUMAN	DNPEP	3687.43	3493.08	3209.65	1507.43	3127.96	2926.18	3550.56	1848.06	3148.42
Q9ULD2	Q9ULD2	MTUS1_HUMAN	MTUS1				1791.02	1551.69	720.068			
Q9ULE6	Q9ULE6	PALD_HUMAN	PALD1							7764	3326.98	8113.32
Q9ULH0	Q9ULH0	KDIS_HUMAN	KIDINS220					2490.5	2116.07	4402.64		2579.23
Q9ULK4	Q9ULK4	MED23_HUMAN	MED23	913.447				868.357	789.587	1175.18	1605.96	656.345
Q9ULL5	Q9ULL5	PRR12_HUMAN	PRR12							1316.21		1433.74
Q9ULT8	Q9ULT8	HECD1_HUMAN	HECTD1					1574.35	2367.41			
Q9ULV0	Q9ULV0	MYO5B_HUMAN	MYO5B							5541.69	6038.55	6992.11
Q9ULV4	Q9ULV4	COR1C_HUMAN	CORO1C	17372.6	14123	15052.6	92478.6	236184	90717.9	103836	60366.8	42766.8
Q9ULW0	Q9ULW0	TPX2_HUMAN	TPX2				1368.94	1987.45	1827.42	1740.1	1198.55	1155.12
Q9ULX3	Q9ULX3	NOB1_HUMAN	NOB1	4828.82	4749.38	3719.82	8652.46	12704.6	10316.3	13886.4	11410.7	13242.7
Q9ULX9	Q9ULX9	MAFF_HUMAN	MAFF	1574.72		544.797	1481.14	1003.76	1018.58	279.595		
Q9UM00	Q9UM00	TMCO1_HUMAN	TMCO1			3165.4	3822.17	4650.35	3898.94	6037.6	2546.77	4747.12
Q9UM54	Q9UM54	MYO6_HUMAN	MYO6		1972.77	1815.24	2681.24	8248.17	5158.72	7589.21	5245.23	7572.17
Q9UMR2	Q9UMR2	DD19B_HUMAN	DDX19B					3571.94	2984.27	2933.89	2048.69	4523.13
Q9UMS4	Q9UMS4	PRP19_HUMAN	PRPF19	10317.2	10425	14434.4	14921.3	22303.9	17017.9	20581.5	11635.4	18868.4
Q9UMX0	Q9UMX0;Q9UHD9	UBQL1_HUMAN	UBQLN1	3499.05		4156.61	1886.18	2076.45	3748.28	2500.35		3365.32
Q9UN37	Q9UN37	VPS4A_HUMAN	VPS4A	19484.7	17501.5	14524.9	28154	7970.72	12881	32719.8	42865.2	25526.3
Q9UN81	Q9UN81	LORF1_HUMAN	L1RE1					1814.45	1025.92	3359.78	3652.71	3312.56
Q9UN86	Q9UN86	G3BP2_HUMAN	G3BP2	4394.1	2886.42	3344.36	3812.37	20239.9	4724.63	4609.24	4332.88	4742.43
Q9UNE7	Q9UNE7	CHIP_HUMAN	STUB1			4280.85	3297.9	4568.22	2362.85	3483.78		3822.2
Q9UNF0	Q9UNF0	PACN2_HUMAN	PACSIN2				8738.58	9436.86	9982.3	5927.02	2656.42	4739.4
Q9UNF1	Q9UNF1	MAGD2_HUMAN	MAGED2	2595.62			3888.48	5135.05	3636.27	6295.66	3822.29	7035.46
Q9UNH7	Q9UNH7	SNX6_HUMAN	SNX6			3325.04	2700.47	4829.02	5773.6	2325.55	2203.14	2392.92
Q9UNL2	Q9UNL2	SSRG_HUMAN	SSR3	13096.7	11822.3	19213.3	4141.52	2652.45	3100.45	8173.37		10710.8
Q9UNM6	Q9UNM6	PSD13_HUMAN	PSMD13	5096.22	8689.86	11623.4	12476.5	12457.5	14965	12313.8	8140.2	10104.2
Q9UNN5	Q9UNN5	FAF1_HUMAN	FAF1				2844.38		2209.47	3241.72	2654.49	2588.91
Q9UNN8	Q9UNN8	EPCR_HUMAN	PROCR	4494.22			1920.52	8968.37	5644.76	3371.87	1658.26	3000.1
Q9UNQ2	Q9UNQ2	DIM1_HUMAN	DIMT1				1769.51	2821.06	1686.86	1489.57	2399.46	1485.36
Q9UNS1	Q9UNS1	TIM_HUMAN	TIMELESS				1346.56	4308.54	1180.61	1506.5	1413.63	1072.42
Q9UNS2	Q9UNS2	CSN3_HUMAN	COPS3						1576.25	1060.04		402.269
Q9UNW1	Q9UNW1	MINP1_HUMAN	MINPP1					2764.08	3202.99	2801.51		2579.55
Q9UNY4	Q9UNY4	TTF2_HUMAN	TTF2				1642.71	4367.46	2422.56	2748.1	2327.12	3290.82
Q9UNZ2	Q9UNZ2	NSF1C_HUMAN	NSFL1C				4356.73		1854.97	2736.65	4524.81	2745.66
Q9UNZ5	Q9UNZ5	L10K_HUMAN	C19orf53					371.687				
Q9UPN3	Q9UPN3	MACF1_HUMAN	MACF1	11234.1	4186.59	7238.17	7501.93	10499.9	8052.41	9543.29	6881.17	8821.11
Q9UPN4	Q9UPN4	CP131_HUMAN	CEP131	6588.13	7060.17	7595.11	3274.42	4748.55	3693.42	3296.57	3268.78	3952.38
Q9UPN7	Q9UPN7	PP6R1_HUMAN	PPP6R1				2723.06		2706.84	2546.12	2148.48	2705.14
Q9UPN9	Q9UPN9	TRI33_HUMAN	TRIM33				1614.93	2177.9	765.63	2891.04	1695.22	1280.47
Q9UPT5	Q9UPT5	EXOC7_HUMAN	EXOC7			1606.15	4662.27	2327.95	2201.42	2415.71	2900.92	2634.61
Q9UPU5	Q9UPU5	UBP24_HUMAN	USP24				1961.42	1915.82	1546.78	1377.21	1882.36	1760.24
Q9UPW6	Q01826;Q9UPW6	SATB2_HUMAN	SATB2					10990.4	3075.24			
Q9UQ03	Q9UQ03	COR2B_HUMAN	CORO2B	26396.5	43975.4	22997.1		7006.95	4489.13	2792.98		
Q9UQ13	Q9UQ13	SHOC2_HUMAN	SHOC2				2906.41			2289.42	2392.25	3232.83
Q9UQ49	Q9UQ49	NEUR3_HUMAN	NEU3					1076.09	1661.78	5229.18		6461.11
Q9UQ80	Q9UQ80	PA2G4_HUMAN	PA2G4	6623.86	3595.54	6517.49	3282.07	10202.5	18858.3	11679.8	3837.48	11444
Q9UQB8	Q9UQB8	BAIP2_HUMAN	BAIAP2				1507.66	2862.15	2368.49	1519.38	618.265	1217.95
Q9UQC2	Q9UQC2	GAB2_HUMAN	GAB2				2812.79	2674.97	1536.23	1368.75	1313.94	1725.94
Q9UQE7	Q9UQE7	SMC3_HUMAN	SMC3				3118.03	3886.08	3982.69	4703.68	4695.79	4340.19
Q9UQR0	Q9UQR0	SCML2_HUMAN	SCML2									
Q9UQR1	Q9UQR1	ZN148_HUMAN	ZNF148					1583.1				

Q 9Y222	Q9Y222	DMTF1_HUMAN	DMTF1		1309.44			1070.89				
Q 9Y224	Q9Y224	RTRAF_HUMAN	RTRAF	5469.86	3679.63	1239.67	3113.26	14326.6	5848.33	4610.03	1865.82	4716.25
Q 9Y230	Q9Y230	RUVB2_HUMAN	RUVBL2	16208.8	13670.2	17158	23733.7	22685	23214.3	13179.2	11169.2	14372.2
Q 9Y237	Q9Y237	PIN4_HUMAN	PIN4					310.058	970.04	395.387		271.701
Q 9Y261	Q9Y261	FOXA2_HUMAN	FOXA2								2629.49	
Q 9Y262	Q9Y262	EIF3L_HUMAN	EIF3L	5455.75	2287.11	4529.49	9156.32	5336.94	6884.66	3541.59	4648.64	4877.93
Q 9Y263	Q9Y263	PLAP_HUMAN	PLAA	6548.79	5481.55	4576.76	5852.96	1593.51	2657.88	1911.61	4720.68	2374.49
Q 9Y265	Q9Y265	RUVB1_HUMAN	RUVBL1	32692.8	28722.6	30879.4	48235.7	48315.9	43074.3	33733.8	29845.8	28620.9
Q 9Y266	Q9Y266	NUDC_HUMAN	NUDC	7890.18	4271.83	6809.43	7787.57	6772.12	14873.7	16970.7	11920	16883
Q 9Y277	Q9Y277	VDAC3_HUMAN	VDAC3	13357.3	20488.8	14232.3	1575.51	8152.79	4917.3	5703.93	5122.32	6633.02
Q 9Y281	Q9Y281	COF2_HUMAN	CFL2			12783.8	1870.14	15485.9	9330.46			
Q 9Y285	Q9Y285	SYFA_HUMAN	FARSA			4036.44	3860.03	3610.18	3174.58	3889.09	3086.78	4002.71
Q 9Y289	Q9Y289	SC5A6_HUMAN	SLC5A6	1807.63			2694.53		3748.9	5840.6	3833.34	4784.4
Q 9Y295	Q9Y295	DRG1_HUMAN	DRG1	4363.58	2616.83	3348.3	2269.73	4584.42	6003.38	5218.39	1208.39	5932.25
Q 9Y2A7	Q9Y2A7	NCKP1_HUMAN	NCKAP1	7220.96	7393.52	12281	5828.48	7048.1	7855.77	5088.61	3290.53	5478.95
Q 9Y2B0	Q9Y2B0	CNPY2_HUMAN	CNPY2			893.39	4009.07		1178.43		3362.59	1135.65
Q 9Y2D4	Q9Y2D4	EXC6B_HUMAN	EXOC6B	2851.02	2898.53	3395.96	1839.27	3218.15	3714.85	1603.41	1229.32	
Q 9Y2D5	Q9Y2D5	AKAP2_HUMAN	AKAP2	5840.32	3152.58	6221.69	3859.14	3624.46	5782.35		1568.27	
Q 9Y2G0	Q9Y2G0	EFR3B_HUMAN	EFR3B							1699.75		1496.39
Q 9Y2H1	Q9Y2H1	ST38L_HUMAN	STK38L			36237.4	6466.47		1626.9	7890.27	11864.7	9493.58
Q 9Y2I6	Q9Y2I6	NINL_HUMAN	NINL		16368.1	8602.21	21062.9	41051.5	28214.7	18957.5	12858.7	17067.3
Q 9Y2J2	Q9Y2J2	E41L3_HUMAN	EPB41L3				3138.09	2371.61	3478.72	1477.43		1274.69
Q 9Y2K3	Q9Y2K3	MYH15_HUMAN	MYH15	20232.6	19024.2	13011.2	77631.6	76174.3	51171	9342.63		8671.16
Q 9Y2K7	Q9Y2K7	KDM2A_HUMAN	KDM2A				2771.64			4489.8		2285.5
Q 9Y2L1	Q9Y2L1	RRP44_HUMAN	DIS3				916.9	2097.42	2793.1	2561.4	1485.57	1991.79
Q 9Y2Q 3	Q9Y2Q3	GSTK1_HUMAN	GSTK1	6781.97	3222.24	4520.07	2019.85	1729.9	2836.49	5276.42	4539.63	3919.87
Q 9Y2Q 5	Q9Y2Q5	LTOR2_HUMAN	LAMTOR2					2093.41	1046.98	1830.82		563.203
Q 9Y2R9	Q9Y2R9	RT07_HUMAN	MRPS7	6300.71	9080.21	7298.93	3928.55	6012.69	3617.69	8442.23	5284.09	9710.43
Q 9Y2S7	Q9Y2S7	PDIP2_HUMAN	POLDIP2	2308.99			3925.54	3256.44	4495.12	3668.92	3254.79	4732.68
Q 9Y2T2	Q9Y2T2	AP3M1_HUMAN	AP3M1	9613.61	9702.19	5531.18	5843	12168.1	7481.26	7254.46	7210.06	8455.05
Q 9Y2U5	Q9Y2U5	M3K2_HUMAN	MAP3K2	28585.9	27609.2		13533.2	7056.61	10465.6		11462.3	29152
Q 9Y2V2	Q9Y2V2	CHSP1_HUMAN	CARHSP1			340.101	253.131		1368.7			
Q 9Y2V7	Q9Y2V7	COG6_HUMAN	COG6				1311.85		1752.93	1688.19	1567.34	1999.69
Q 9Y2W1	Q9Y2W1	TR150_HUMAN	THRAP3	2724.23	3400.07			2442.76	1054.57	1093.06		
Q 9Y2W2	Q9Y2W2	WBP11_HUMAN	WBP11					9746.73	8611.56	4753.15		3308.85
Q 9Y2W6	Q9Y2W6	TDRKH_HUMAN	TDRKH						539.065	1335.63	238.963	1398.97
Q 9Y2X0	Q9Y2X0	MED16_HUMAN	MED16	1921.47	4358.37	2943.86	3239.13	2211.12	2249.03	4504.9	3752.91	4689.66
Q 9Y2X3	Q9Y2X3	NOP58_HUMAN	NOP58				1860.79	3186.56	2243.37	1803.45	1258.63	1842.26
Q 9Y2X7	Q9Y2X7	GIT1_HUMAN	GIT1			2420.56	3618.7		859.389	4994.3	2159.68	5506.87
Q 9Y2X9	Q9Y2X9	ZN281_HUMAN	ZNF281					1583.09	384.53			
Q 9Y2Y1	Q9Y2Y1	RPC10_HUMAN	POLR3K									
Q 9Y2Y9	Q9Y2Y9	KLF13_HUMAN	KLF13									
Q 9Y2Z0	Q9Y2Z0	SGT1_HUMAN	SUGT1				1442.55	978.455	2154.56	1597.65		1518.4
Q 9Y2Z2	Q9Y2Z2	MTO1_HUMAN	MTO1	4514.3			1600.6	2732.6	1400.27	3350.61	2908.7	3845.82
Q 9Y305	Q9Y305	ACOT9_HUMAN	ACOT9	11435.4	10912.8	10629.2	12346	8013.95	5598.11	10193.2	17720.8	10973.2
Q 9Y312	Q9Y312	AAR2_HUMAN	AAR2		1479.37			2152.37	1056.21	2051.05	2911.42	3442.54
Q 9Y314	Q9Y314	NOSIP_HUMAN	NOSIP					3108.31	2792.14	944.047		1505.94
Q 9Y315	Q9Y315	DEOC_HUMAN	DERA		3282.74	4386.15	4639.05	2736.5	3938.76	6292.66	4977.68	5614.28
Q 9Y320	Q9Y320	TMX2_HUMAN	TMX2		2013.89		1787.2	4464.05	4649.97	7142.75	2427.46	3511.37
Q 9Y371	Q9Y371	SHLB1_HUMAN	SH3GLB1				1976.87	2625.76	3271.04	1805.07		2424.3
Q 9Y383	Q9Y383	LC7L2_HUMAN	LUC7L2				1941.62	1832.52	1180.23		2501.74	1620.77
Q 9Y3B3	Q9Y3B3	TMED7_HUMAN	TMED7				3893.55	6685.2	10558.1	9019.2	3603.45	8140.94
Q 9Y3B4	Q9Y3B4	SF3B6_HUMAN	SF3B6	1499.74	2805.49			1253.44	1045.49		1574.8	917.339
Q 9Y3B7	Q9Y3B7	RM11_HUMAN	MRPL11	6234.1	7113.32	5935.23	3289.71	6177.46	3655.84	3641.89	6703.93	3380.07
Q 9Y3C1	Q9Y3C1	NOP16_HUMAN	NOP16			4586.3		22743.4	26777.4			
Q 9Y3C6	Q9Y3C6	PPIL1_HUMAN	PPIL1			11773.9	2161.8	2099.68	2735.04			2216.14
Q 9Y3D0	Q9Y3D0	CIA2B_HUMAN	CIAO2B				1578.22	1256.53	2813.93	2930.61	2134.12	4242.52
Q 9Y3D3	Q9Y3D3	RT16_HUMAN	MRPS16	1568.05	4280.33	1555.32	1005.38	1130.64	1419.17	2276.95	1304.56	2937.66
Q 9Y3D6	Q9Y3D6	FIS1_HUMAN	FIS1					1866.2	1026.96	1956.18	1262.8	2518.72
Q 9Y3D8	Q9Y3D8	KAD6_HUMAN	AK6					2839.57	1668.62	2125.36		3698.28
Q 9Y3D9	Q9Y3D9	RT23_HUMAN	MRPS23	7047.49		1948.76	3899.9	2268.91	2759.99	4180.69	5851.94	4152.03
Q 9Y3E5	Q9Y3E5	PTH2_HUMAN	PTRH2		800.468			2331.74	1728.02	823.142		1043.39
Q 9Y3F4	Q9Y3F4	STRAP_HUMAN	STRAP	1285.12		2101.82	2088.1	2787.01	6302.54	4140.27	1282.39	3758.92
Q 9Y3I0	Q9Y3I0	RTCB_HUMAN	RTCB	8943.79	3561.84	5831.44	8719.69	15102.9	10090.4	9293.14	7702.14	9708.99
Q 9Y3I1	Q9Y3I1	FBX7_HUMAN	FBXO7				2695.01	2017.56	1868.88	1587.53	1909.76	2535.13
Q 9Y3L5	Q9Y3L5	RAP2C_HUMAN	RAP2C				3851.94	3858.46	4953.45	14987.7	3706.16	14296.2
Q 9Y3M8	Q9Y3M8	STA13_HUMAN	STARD13	2310.26	1312.76	915.518	542.682	594.641	632.68			120.595
Q 9Y3R4	Q9Y3R4	NEUR2_HUMAN	NEU2									
Q 9Y3R5	Q9Y3R5	DOP2_HUMAN	DOP1B				593.452	1951.53	1711.02	2027.09		637.394
Q 9Y3T6	Q9Y3T6	R3HC1_HUMAN	R3HCC1				5739.24	3228.21	4816.43	4427.81	5963.51	4393.46
Q 9Y3U8	Q9Y3U8	RL36_HUMAN	RPL36	12027.6	17504.1	22125.7	24563.7	45466.6	17364.6	24407.5	24897	25236.8
Q 9Y3Z3	Q9Y3Z3	SAMH1_HUMAN	SAMHD1	24260.2	21938.4	18536.3	14488.9	11579.8	10395.6	13962.4	25554.1	18648.7
Q 9Y421	Q9Y421	FA32A_HUMAN	FAM32A				3128.09	3576.62	2996.96		2572.54	2628.47
Q 9Y446	Q9Y446	PKP3_HUMAN	PKP3				6247.81	10626.8	4978.52	8096.58	8097.72	8985.41
Q 9Y467	Q9Y467	SALL2_HUMAN	SALL2					441.493	335.667			
Q 9Y487	Q9Y487	VPP2_HUMAN	ATP6V0A2					5245.22	1966.85	2639.28		1488.86
Q 9Y490	Q9Y490	TLN1_HUMAN	TLN1	13751.4	11348.5	11526.7	13811.7	10548.8	12570	6707.58	5482.43	7221.61
Q 9Y4B5	Q9Y4B5	MTCL1_HUMAN	MTCL1				4613.06	6145.49	2960.31	1055.94	1295.08	1227.86
Q 9Y4C2	Q9Y4C2	TCAF1_HUMAN	TCAF1			3307.99	2852.86	5123.37	3130.15	5319.74	6517.24	8256.6
Q 9Y4D1	Q9Y4D1	DAAM1_HUMAN	DAAM1			3495.16	1745.25	3851.99	3041.07	3361.63	1491.82	2359.61
Q 9Y4F1	Q9Y4F1	FARP1_HUMAN	FARP1	9762.59	6198.98	5587.4	2061.86		3936.93	14556.5	5327.33	16003.1
Q 9Y4F3	Q9Y4F3	MARF1_HUMAN	MARF1					3130.73	1194.65	832.133		
Q 9Y4H2	Q9Y4H2	IRS2_HUMAN	IRS2	11055	8018.8	9236.06	6364.6	5899.15	4555.08	1470.75	1297.47	2027.29
Q 9Y4I1	Q9Y4I1	MYO5A_HUMAN	MYO5A	12207	13103.7		22765.3	74388.2	36884.1	38025.6	23844.7	35136.2
Q 9Y4K0	Q9Y4K0	LOXL2_HUMAN	LOXL2					6159.81	5089.15			
Q 9Y4L1	Q9Y4L1	HYOU1_HUMAN	HYOU1	5167.94	2536.06	5848.74	9065.3	9491.88	14428.2	7728.3	3955.16	7849.47
Q 9Y4P3	Q9Y4P3	TBL2_HUMAN	TBL2							1856.37	1918.58	1345.2
Q 9Y4R8	Q9Y4R8	TELO2_HUMAN	TELO2	1335.58	1029.78	975.307	7001.4	4979.54	4409.37	4682.33	4984.95	6524.85
Q 9Y4W6	Q9Y4W6	AFG32_HUMAN	AFG3L2				2402.11	2483.52	3881.95	3967.75	3281.22	5804.1
Q 9Y4X0	Q9Y4X0	AMMR1_HUMAN	AMMECR1					2343.92	2018.96	2346.97		2436.3
Q 9Y4X4	Q9Y4X4	KLF12_HUMAN	KLF12									
Q 9Y4Z0	Q9Y4Z0	LSM4_HUMAN	LSM 4.00					1782.36	2082.04	1282.49		1526.3
Q 9Y512	Q9Y512	SAM50_HUMAN	SAMM50	42610.5	38171.9	32078.3		11210.2	4988.79	10852.4		10234.1
Q 9Y520	Q9Y520	PRC2C_HUMAN	PRRC2C	2595.11	1468.8	1874.5	2539.93	5123.59	3231.54	2266.85	1709.48	1892.09
Q 9Y570	Q9Y570	PPME1_HUMAN	PPME1				1539.06	2426.76	3438.94	1369.52		
Q 9Y5A7	Q9Y5A7	NUB1_HUMAN	NUB1			1006.47	1221.54	1012.22	866.119	1688.85	787.494	1622.49
Q 9Y5A9	Q9Y5A9	YTHD2_HUMAN	YTHDF2					10041.3	4951.46		2936.75	6196.53
Q 9Y5B9	Q9Y5B9	SP16H_HUMAN	SUPT16H	7688.07	7586.79	4196.91	6899.18	7934.53	4644.4	5828.99	5102.44	6281.67
Q 9Y5K5	Q9Y5K5	UCHL5_HUMAN	UCHL5				3166.35	5075.76	5196.39	2370.42	2512.56	2108.18
Q 9Y5K6	Q9Y5K6	CD2AP_HUMAN	CD2AP				4405.32	4781.6	8327.58	2621.36	2138.5	3433.91
Q 9Y5K8	Q9Y5K8	VATD_HUMAN	ATP6V1D	3260.21	4357.03	5806.96	2461.91	5847.89	5601.81	3870.87		4006.76
Q 9Y5L0	Q9Y5L0	TNPO3_HUMAN	TNPO3	5643.74	5781.85	7254.99	7307.33	9049.16	3905.26	10109.4	8541.25	11174.7
Q 9Y5L4	Q9Y5L4	TIM13_HUMAN	TIMM13	3539.95	2215.28	2285.56	5824.61	9249.12	12656	10753.2	5364.76	8332.6
Q 9Y5M8	Q9Y5M8	SRPRB_HUMAN	SRPRB	2630.85	2860.65	3431.65	7030.48	4266.21	2917.45	7493.56	8484.68	6348.03
Q 9Y5P6	Q9Y5P6	GMPPB_HUMAN	GMPPB				3247.25				4641.69	
Q 9Y5Q 8	Q9Y5Q8	TFIC5_HUMAN	GTF3C5	2839.9			2396.76	2191.21	3172.07	4027.04	3318.54	4658.96
Q 9Y5Q 9	Q9Y5Q9	TFIC3_HUMAN	GTF3C3		2514.01		1610.08	2373.19	2613.34	3964.44	1815.91	3562.61
Q 9Y5S1	Q9Y5S1	TRPV2_HUMAN	TRPV2				5280.08	5487.44	4485.99	3078.04		4664.61
Q 9Y5S2	Q9Y5S2;Q5VT25	MRCKB_HUMAN	CDC42BPB	4847.7	6992.82	3500.88	3185.18	3510.85	3582.57	4279.08	2506.56	4458.2
Q 9Y5V3	Q9Y5V3	MAGD1_HUMAN	MAGED1	5394.87	4328.76	5491.35	3948.93	3066.4	2084.77	5433.26	9866.1	5428.84
Q 9Y5X1	Q9Y5X1	SNX9_HUMAN	SNX9	6949.96		6117.23	4555.8	3330.73	5008.74	2236.33		3415.18
Q 9Y5X2	Q9Y5X2	SNX8_HUMAN	SNX8	3900.85	3877.98	4161.23	4205.45	963.974	1942.37	2090.42	2185.92	2441.48
Q 9Y5X3	Q9Y5X3	SNX5_HUMAN	SNX5						2117.08	1013.6		591.94
Q 9Y5Y0	Q9Y5Y0	FLVC1_HUMAN	FLVCR1					760.36	843.826	2518.24		1605.85
Q 9Y605	Q9Y605	MOFA1_HUMAN	MRFAP1		1904.01	1270.34	546.726	1211.35	2309.29	1261.86		1684.6
Q 9Y606	Q9Y606	PUS1_HUMAN	PUS1			1708.01	2691.67		4086.14	5149.34	2475.42	5718.24
Q 9Y613	Q9Y613	FHOD1_HUMAN	FHOD1				2322.43	6635.03	4919.48	2273.51	2457.63	2068.97
Q 9Y617	Q9Y617	SERC_HUMAN	PSAT1	7165.64	4404.86	5659.77	10977.9	8106.15	16824.6	6219.96	4134.23	4675.23
Q 9Y618	Q9Y618	NCOR2_HUMAN	NCOR2			2207.19	1629.59	3615.15	3943.71	3494.45	1522.26	4756.3
Q 9Y624	Q9Y624	JAM1_HUMAN	F11R				2411.05	4858.04	4265.44	12113.4		9789.83
Q 9Y639	Q9Y639	NPTN_HUMAN	NPTN	4336.58	4330.74	5344.54	3445.68	7652.43	5894.82	2733.48		2396.99
Q 9Y666	Q9Y666	S12A7_HUMAN	SLC12A7							3103.98		3910.29
Q 9Y673	Q9Y673	ALG5_HUMAN	ALG5				3813.07				4101.68	
Q 9Y678	Q9Y678	COPG1_HUMAN	COPG1	9848.11	8040.86	10173.4	14109.9	9359.75	8234.1	9619.59	9299.48	9999.04
Q 9Y679	Q9Y679	AUP1_HUMAN	AUP1	1423.18	1781.52	1428.37	3301.65	3369.47	4385.25	7517.77	1777.87	8013.77
Q 9Y696	Q9Y696	CLIC4_HUMAN	CLIC4	3792.44	3326.09	4914.65	2906.13		2062.41			
Q 9Y697	Q9Y697	NFS1_HUMAN	NFS1			2452.88	2488.88	2652.46	2153.98	2744.23	3942.03	3779.65
Q 9Y6C9	Q9Y6C9	MTCH2_HUMAN	MTCH2		11592	12495.1		5119.58	3569.55	8249.47	3175.69	7830.89
Q 9Y6D5	Q9Y6D6;Q9Y6D5	BIG2_HUMAN	ARFGEF2	1545.14			4797.14	4520.7	3787.31	3974.47	5132.93	3969.94
Q 9Y6D6	Q9Y6D6	BIG1_HUMAN	ARFGEF1	3134.44	4240.44	2559.52	3737.56	2389.08	3079.34	3366.39	2673.54	3956.51
Q 9Y6E0	O00506;Q9Y6E0	STK24_HUMAN	STK24	1928.11	1519.32	2097.96	1568.11	8951.96	3479.92	3215.27	1216.42	3107.89
Q 9Y6E2	Q9Y6E2	5MP1_HUMAN	BZW2	6109.25	4211.44	3611.5	7218.49	6166.45	7899.03	9246.45	4615.23	8711.47
Q 9Y6G9	Q9Y6G9	DC1L1_HUMAN	DYNC1LI1	4247.42	6253.87	8098.47	10680.3	7540.7	6344.05	6827.38	9644.69	7621.88
Q 9Y6J9	Q9Y6J9	TAF6L_HUMAN	TAF6L	4816.23	3485.18	3797.1	1854.3	7140.06		6819.49	4282.91	7913.76
Q 9Y6K0	Q9Y6K0	CEPT1_HUMAN	CEPT1	7533.7	7772.49	4925.49	5224.65	5760.63	3812.93	11895.1	7110.08	15948.5
Q 9Y6K5	Q9Y6K5	OAS3_HUMAN	OAS3	4452.85	3324.69	3728.21	2640.99	4421.12	1904.95	2536.63	11576.9	3496.62
Q 9Y6K8	Q9Y6K8	KAD5_HUMAN	AK5	5301.86	2416.8	5796.63						
Q 9Y6M1	Q9Y6M1	IF2B2_HUMAN	IGF2BP2	9441.43	6780.44	7037.14	4915.4	8831.42	3929.85	2976.58	3321.82	2558.44
Q 9Y6M4	Q9Y6M4;P78368;Q9HCP0	KC1G3_HUMAN	CSNK1G3				6312.54	12541.8	11647.8	38637.1	9624.57	29625.9
Q 9Y6M7	Q9Y6M7	S4A7_HUMAN	SLC4A7					1049.67	414.339	1065.33		1869.15
Q 9Y6N5	Q9Y6N5	SQOR_HUMAN	SQOR	41651.9	31822.8	42817				11606.8	16047.3	14247.7
Q 9Y6N7	Q9Y6N7	ROBO1_HUMAN	ROBO1	1408.06		1096.86		973.143	1178.54	2372.51	1203.91	2092.39
Q 9Y6Q 1	Q9Y6Q1	CAN6_HUMAN	CAPN6							2382.61		2206.29
Q 9Y6Q 9	Q9Y6Q9	NCOA3_HUMAN	NCOA3									
Q 9Y6R9	Q9Y6R9	CCD61_HUMAN	CCDC61	10653.8	9698.03	9733.35			2092.17			
Q 9Y6V7	Q9Y6V7	DDX49_HUMAN	DDX49				1599.43	5535.06	4363.02	3355.87	4511.82	3484.61
Q 9Y6W5	Q9Y6W5	WASF2_HUMAN	WASF2				4532.58	1531.43	2581.08	614.923		923.282
Q 9Y6X9	Q9Y6X9	MORC2_HUMAN	MORC2				686.516	675.432		1065.59	276.819	1092.7
Q 9Y6Y8	Q9Y6Y8	S23IP_HUMAN	SEC23IP	4454.89		3227.22	3952.03	10427.3	11092.1	6359.16	1739.71	5245.94