

-Log ANOVA	ANOVA q	Protein IDs	Majority p	Protein na	Gene names
0,741374	0,193349	A0A024QZ	A0A024QZ	Serpin B6	SERPINB6
1,63684	0,041581	E7EMB6;A	E7EMB6;A	Aspartyl ar	DNPEP
3,58429	0,002797	A0A024RA	A0A024RA	Proteasom	PSMA2
1,22098	0,082771	A0A087X0	A0A087X0	Ig kappa c	IGKV2-40;IGKV2D-28;IGKV A18;IGKV2D-26;IC
3,94102	0,001722	A0A075B7	A0A075B7	Epiplakin	EPPK1
1,73782	0,035297	Q16881;A	Q16881;A	Thioredoxi	TXNRD1
4,8719	0,000587	A0A087X2	A0A087X2	Glutathion	GPX4
8,14716	0	A0A087W1	A0A087W1	Collagen al	COL1A2
1,32598	0,069517	P11940;A0	P11940;A0	Polyadenyl	PABPC1
3,66879	0,002533	J3QTS8;A0	J3QTS8;A0	Tryptase al	TPSAB1;TPSB2
4,46716	0,000936	A0A087W1	A0A087W1	26S protea	PSMD13
5,61989	0,000321	A0A087W1	A0A087W1	Glutathion	GPX1
9,39158	0	A0A087W1	A0A087W1	MICOS con	APOOL
13,9745	0	A0A087W1	A0A087W1	Aromatic-L	DDC
4,67834	0,000721	A0A087W1	A0A087W1	Clathrin he	CLTC
3,02143	0,005617	A0A087W1	A0A087W1	Ribosome-	RRBP1
5,56058	0,000357	H0YMB3;H	H0YMB3;H	GMP reduc	GMPR2
1,01064	0,119382	P06753-2;I	P06753-2;I	Tropomyo:	TPM3;DKFZp686J1372
7,48863	2,78E-05	A0A087W1	A0A087W1	NADH deh	NDUFA10
1,12196	0,097776	A0A087W1	A0A087W1	Basal cell a	BCAM
0,611613	0,246917	A0A087W1	A0A087W1	AP-2 comp	AP2M1
2,7622	0,007883	A0A087W1	A0A087W1	Tumor pro	TPD52L2
1,09672	0,10223	A0A087W1	A0A087W1	UTP--gluco	UGP2
2,96653	0,006034	A0A087W1	A0A087W1	Prostaglan	PTGES3
2,61753	0,009509	A0A087W1	A0A087W1	Lipoma-pr	LPP
17,7011	0	A0A087W1	A0A087W1	Isocitrate c	IDH3B
4,2845	0,001157	A0A087W1	A0A087W1	Methyltrar	METTL7B
5,28093	0,000445	A0A087X0	A0A087X0	Hypoxia u	HYOU1
8,93732	0	P09668;A0	P09668;A0	Pro-cathep	CTSH
7,61547	1,92E-05	A0A087X0	A0A087X0	Collagen al	COL15A1
11,3335	0	A0A087X0	A0A087X0	Collagen al	COL6A1
10,6351	0	A0A087X0	A0A087X0	Acyl-coenz	ACOT2
6,61502	8,59E-05	A0A087X1	A0A087X1	Septin-8	SEPT8
5,54128	0,000375	A0A087X1	A0A087X1	Ras-relate	RAB18
10,0058	0	A0A087X1	A0A087X1	Glutathion	GPX3
2,8707	0,006922	A0A087X1	A0A087X1	Proteasom	PSME2
2,89992	0,006684	A0A087X2	A0A087X2	Basigin	BSG
0,506614	0,303523	A0A087X2	A0A087X2	26S protea	PSMC6
9,56058	0	PODPI2;A0	PODPI2;A0		C21orf33
13,0383	0	A0A0A0M1	A0A0A0M1	Laminin su	LAMA4
4,12626	0,001421	F2Z357;A0	F2Z357;A0	Rap1 GTPa	RAP1GAP
2,58299	0,009985	A0A0A0M1	A0A0A0M1	Heterogen	HNRNPUL1
8,0359	1,04E-05	G3V5X4;A	G3V5X4;A	Nesprin-2	SYNE2
4,9694	0,000533	A0A0A0M1	A0A0A0M1	Unconvent	MYO6
3,76803	0,002262	P01857;A0	P01857;A0	Ig gamma-	IGHG1
1,2248	0,082189	A0A0A0M1	A0A0A0M1	Sideroflexi	SFXN3
1,54916	0,048195	A0A0A0M1	A0A0A0M1	Plastin-3	PLS3
14,1118	0	A0A0A0M1	A0A0A0M1	Aldo-keto	AKR1C3
3,75115	0,002302	A0A0G2JK	A0A0G2JK	Tapasin	TAPBP

0,939432 0,134572 J3KR24;A0. J3KR24;A0. Isoleucine- IARS
8,21323 0 A0A0A0M1 A0A0A0M1 Isovaleryl- (IVD
6,50634 0,000106 A0A0A0M1 A0A0A0M1 E3 ubiquitin RNF213
3,8442 0,002014 A0A0A0M1 A0A0A0M1 Integrin-like ILK
8,47581 0 A0A0R4J2C A0A0R4J2C Neutral chain NCEH1
6,70839 7,1E-05 A0A1B0GV A0A1B0GV Adenylosin ADSL
4,7136 0,000686 A0A0A6YY. A0A0A6YY. Transmembrane TMED7-TICAM2;TMED7
2,51132 0,011041 A0A0A6YY. A0A0A6YY. UPF0562 protein LUC7L2;C7orf55
0,811789 0,170062 A0A0A6YY| A0A0A6YY| Coronin;Cc CORO7-PAM16;CORO7
8,82781 0 A0A0B4J1F A0A0B4J1F Acyl-CoA synthase ACS3
6,58397 8,53E-05 A0A0B4J1F A0A0B4J1F 4-hydroxyphenyl HPD
1,72397 0,036092 A0A0B4J1Z A0A0B4J1Z Serine/arginine SRSF7
1,35596 0,066096 A0A0B4J2E A0A0B4J2E Immunoglobulin IGLL5;IGLC1
8,11711 0 A0A0B4J2E A0A0B4J2E Ig heavy chain IGHV3OR16-9;IGHV3OR16-10;IGHV3-20;IGHV
0,789304 0,177323 Q5W0H4;A Q5W0H4;A Translatin TPT1
3,72996 0,002364 G3V5T0;AC G3V5T0;AC Maleylacetate GSTZ1
1,08644 0,104146 A0A0C4DF A0A0C4DF Transmembrane TM9SF4
3,2526 0,004198 A0A0C4DF A0A0C4DF Monoglyceride MGLL
0,682999 0,214984 A0A0C4DF A0A0C4DF Protein SET SET;SETSIP
4,22218 0,001241 C9J9K3;A0. C9J9K3;A0. 40S ribosome RPSA
2,18051 0,017601 A0A0C4DG A0A0C4DG Serine protein HTRA2
22,1495 0 A0A0C4DG A0A0C4DG Enoyl-CoA hydratase ECI2
7,34631 2,71E-05 A0A0C4DG A0A0C4DG Calpain small CAPNS1
6,36573 0,000138 A0A0C4DG A0A0C4DG Dolichyl-diphosphate DDOST
3,3784 0,003574 A0A0C4DH A0A0C4DH IGHV5-51
5,30515 0,000442 Q30118;AC Q30118;AC HLA class II HLA-DRA
2,18395 0,017473 H9KV28;AC H9KV28;AC Protein disulfide DIAPH1
0,223456 0,551644 A0A0G2JIV A0A0G2JIV Heat shock HSPA1B;HSPA1A
4,00668 0,001625 A2ACR1;AC A2ACR1;AC Proteasome PSMB9
1,52627 0,049862 Q5SRQ3;Q Q5SRQ3;Q Casein kinase CSNK2B-LY6G5B-1181;CSNK2B;CSNK2B-LY6C
3,38051 0,003578 A0A0G2JIM A0A0G2JIM Ig alpha-2 (IGHA2
1,0295 0,115685 A0A0G2JPI A0A0G2JPI
0,670886 0,219859 O94973;AC O94973;AC AP-2 component AP2A2
1,13349 0,095703 A0A0J9YYE A0A0J9YYE Ig heavy chain IGHV3-23
2,70238 0,008529 A0A0J9YYC A0A0J9YYC Serum protein PON2
3,83431 0,002033 P43243;A8 P43243;A8 Matrin-3 MATR3
5,29364 0,000441 A0A0U1RC A0A0U1RC Fatty acid synthase FASN
4,02863 0,001583 A0A0U1RR A0A0U1RR Spermatogonein SPATA20
1,51519 0,050732 A0A140T9I A0A140T9I HLA class I HLA-A
3,16094 0,004669 Q03518;AC Q03518;AC Antigen peptide TAP1
9,82734 0 A0A140TA A0A140TA Protein QIL QIL1
7,17488 4,31E-05 A0A182DV A0A182DV Thioredoxin TXNRD2
7,06457 5,07E-05 H0Y695;B3 H0Y695;B3 Aminomethylase AMT
6,51949 0,000107 Q13510-2; Q13510-2; Acid ceramidase ASAH1
3,13551 0,004826 A0A1B0GL A0A1B0GU Ig mu chain IGHM
4,69619 0,000706 A0A1B0GV A0A1B0GV Kinesin-like KIF21A
15,2434 0 A0A1B0GV A0A1B0GV Cathepsin B CTSD
15,3123 0 A0A1C7CY A0A1C7CY Dihydropyridin DPYSL2
1,36818 0,064734 A0A1W2PI A0A1W2PI
3,86397 0,001967 A0A1W2PC A0A1W2PC NAD-dependent ME2

3,61948 0,002677 A0A1W2PF A0A1W2PF Lysosome | SCARB2
1,49833 0,052066 A0A1X7SB; A0A1X7SB; Probable A DDX17
2,06897 0,021087 A0A286YE; A0A286YE; Ig gamma- IGHG3
2,87343 0,006896 A0A286YE; A0A286YE; Ig alpha-1 | IGHA1
3,43343 0,003425 A0A286YE; A0A286YE; Ig gamma- IGHG2
4,58555 0,000823 A0A286YF; A0A286YF; D-3-phospho PHGDH
3,10596 0,005043 A0A2C9F2I A0A2C9F2I Palmitoyl- f PPT1
1,52747 0,049809 A0A286YFJ A0A286YFJ Ig gamma- IGHG4
2,67996 0,008785 A0A2Q2TT A0A2Q2TT Ig kappa c f
1,10949 0,099982 A0A2R8Y6I A0A2R8Y6I Probable h HARS2
5,4671 0,000394 O60313-1C O60313-1C Dynamin-li OPA1
14,9195 0 A6NNI4; AC A6NNI4; AC Tetraspani CD9
4,16099 0,001344 A0A2R8Y4; A0A2R8Y4; Tensin-1 TNS1
4,5519 0,000844 P35222; A0 P35222; A0 Catenin be CTNNB1
0,465629 0,328074 A0A2R8Y6. A0A2R8Y6. 60S riboso RPL5
3,85412 0,001998 A0A2R8YD A0A2R8YD Succinyl-Cc SUCLA2
4,73637 0,000666 A0A2R8YD A0A2R8YD ATP-depen DDX3X; DDX3Y
1,53859 0,048855 A0A2R8Y7; A0A2R8Y7; Hemoglobi HBG1
16,0428 0 P08237-3; I P08237-3; I ATP-depen PFKM
5,03081 0,000515 A0A2R8YD A0A2R8YD Peroxisom HSD17B4
12,136 0 A0A2R8YG A0A2R8YG Tripeptidyl TPP1
7,47092 2,76E-05 E9PNW4; A E9PNW4; A CD59 glycc CD59
4,32392 0,00112 V9GYY3; AC V9GYY3; AC C-1-tetrahy MTHFD1
12,6258 0 A0A3B3IS8 A0A3B3IS8 Fructose-b ALDOB
2,64994 0,009088 A0A3B3ISK A0A3B3ISK Aspartate- DARS2
2,01918 0,022753 A0A3B3ISC A0A3B3ISC Epidermal EPS8L2
7,01483 5,87E-05 A0A3B3ISS A0A3B3ISS Transmem GPNMB
6,64465 7,05E-05 A0A3B3ISv A0A3B3ISv Collagen al COL4A1
3,06949 0,005332 A0A3B3ISX A0A3B3ISX Tenascin-X TNXB
2,8991 0,006685 E9PG39; AC E9PG39; AC Acylglycerc AGK
3,03955 0,005523 A0A494C0; A0A494C0; Tripeptidyl TPP2
1,07608 0,105939 A0A494C1; A0A494C1; Lipopolysa LRBA
0,923548 0,138176 M0R026; A M0R026; A Acetolacta ILVBL
4,98939 0,000529 A3KMH1; A A3KMH1; A von Willeb VWA8
4,06479 0,001529 A6NJA2; P5 A6NJA2; P5 Ubiquitin c USP14
5,82102 0,000262 E9PJK1; E9I E9PJK1; E9I Tetraspani CD81
0,848422 0,159361 Q99829; BC Q99829; BC Copine-1 CPNE1
5,50566 0,000383 B0QY89; Q; B0QY89; Q; Eukaryotic EIF3L
1,81233 0,031363 B0YIW6; P4 B0YIW6; P4 Coatomer | ARCN1
3,68812 0,002464 B1AJY7; O7 B1AJY7; O7 26S protea PSMD10
7,30325 3,58E-05 B1AKZ5; Q1 B1AKZ5; Q1 Astrocytic | PEA15
4,4826 0,000918 B1ALA9; P6 B1ALA9; P6 Ribose-phc PRPS1
2,06739 0,021095 P07910-2; I P07910-2; I Heterogen HNRNPC
9,95852 0 B3KSI3; O1; B3KSI3; O1; Branched-c BCAT2
18,6177 0 O75390; B4 O75390; B4 Citrate syn CS
7,26409 3,52E-05 B4DLN1; P5 B4DLN1
4,42178 0,001011 B4DUR8; P2 B4DUR8; P2 T-complex CCT3
3,05308 0,005442 B4DXZ6; P5 B4DXZ6; P5 Fragile X m FXR1
4,41086 0,001026 B4DZG7; P2 B4DZG7; P2 ADP-ribosy ARL1
3,58152 0,002816 B4E1Z4; P0 B4E1Z4; P0 Compleme CFB

2,02318 0,022608 B5MC53;B B5MC53;B Protein Mç MPV17
4,56569 0,000837 P16422;B5 P16422;B5 Epithelial c EPCAM
1,18384 0,087867 B5MDF5;P B5MDF5;P GTP-bindir RAN
1,3957 0,061776 Q99613;Bç Q99613;Bç Eukaryotic EIF3C;EIF3CL
3,59791 0,002764 E7EW69;Eç E7EW69;Eç Septin-10 SEPT10
1,3111 0,071058 B7Z6P0;H3 B7Z6P0;H3 Cytosolic F NUBP2
2,06449 0,021153 B7ZKJ8;Q1 B7ZKJ8;Q1 Inter-alpha ITIH4
1,70044 0,037401 C9J0K6;P3ç C9J0K6;P3ç Sorcin SRI
2,71779 0,008383 C9J4Z3;P6ç C9J4Z3;P6ç 60S riboso RPL37A
11,1085 0 C9JAW5;Cç C9JAW5;Cç HIG1 domæ HIGD1A
1,1887 0,08721 C9JD37;Q1 C9JD37;Q1 Growth fac GRB14
0,704628 0,206636 C9JF17;P0ç C9JF17;P0ç Apolipoprc APOD
13,9472 0 P99999;C9 P99999;C9 Cytochrom CYCS
0,926159 0,137775 F5GZZ9;C9 F5GZZ9;C9 Scavenger CD163
7,78092 1,99E-05 C9JIF9;P13 C9JIF9;P13 Acylamino APEH
11,5537 0 C9JIZ6;P07 C9JIZ6;P07 Prosaposin PSAP
3,29645 0,00397 C9JUE0;C9 C9JUE0;C9 Protein TFç TFG
5,78674 0,000267 C9JKI3;Q0ç C9JKI3;Q0ç Caveolin;C CAV1
12,8829 0 C9JRZ6;Q9 C9JRZ6;Q9 MICOS con CHCHD3
4,07657 0,001494 C9JYN0;Q1 C9JYN0;Q1 Synaptoph SYPL1
4,82187 0,000605 O60716;Oç O60716;Oç Catenin de CTNND1
2,4074 0,012859 CON__P00 CON__P00
8,239 0 CON__P02 CON__P02
5,24504 0,000442 CON__P04 CON__P04
6,47604 0,000104 CON__P05 CON__P05
11,3149 0 CON__P05 CON__P05
14,7147 0 CON__P08 CON__P08
4,43779 0,000988 CON__P13 CON__P13
3,41343 0,003493 CON__P35 CON__P35
0,212343 0,56515 CON__P35 CON__P35
10,6197 0 CON__P67 CON__P67
16,1478 0 CON__Q3k CON__Q3k
3,03047 0,005567 D6R934;PC D6R934;PC Compleme C1QB
4,48587 0,000914 Q13557;Qç Q13557;Qç Calcium/cæ CAMK2D
1,05939 0,109253 D6R9H7;Eç D6R9H7;Eç ATP synthæ ATP5G1;ATP5G2;ATP5G3
6,58654 8,54E-05 D6R9P3;Dç D6R9P3;Dç Heterogen HNRNPAB
2,51825 0,010956 D6RCD0;Q D6RCD0;Q Estradiol 1 HSD17B11
3,59126 0,002774 D6RD47;Pç D6RD47;Pç 40S riboso RPS23
2,03988 0,0221 D6RGI3;Dç D6RGI3;Dç Septin-11 SEPT11
0,958965 0,130048 O94979-6; O94979-6; Protein tra SEC31A
1,51569 0,050747 D6RF35;PC D6RF35;PC Vitamin D- GC
0,759122 0,187221 D6RG13;Pç D6RG13;Pç 40S riboso RPS3A
0,133591 0,682456 D6RGG3;Q D6RGG3;Q Collagen al COL12A1
0,744328 0,192322 E5RI99;P6ç E5RI99;P6ç 60S riboso RPL30
0,492745 0,311813 Q7L5N1;Eç Q7L5N1;Eç COP9 signæ COPS6
1,99484 0,02358 J3QLD9;E7 J3QLD9;E7 Flotillin-2 FLOT2
3,2762 0,004094 G3V2V8;J3 G3V2V8;J3 Epididymal NPC2
1,67831 0,038783 E7EV99;E7 E7EV99;E7 Alpha-addç ADD1
8,83381 0 E7ES33;E7ç E7ES33;E7ç Septin-7 SEPT7
1,44638 0,05678 E7EQR4;P1 E7EQR4;P1 Ezrin EZR

2,21404 0,01675 E7ER44;P0 E7ER44;P0 Lactotrans LTF
 7,85021 2,01E-05 E7ER45;O4 E7ER45;O4 Maltase-gl MGAM
 3,34859 0,003683 E7EX90;Q1 E7EX90;Q1 Dynactin s DCTN1;DKFZp686E0752
 1,23377 0,081058 E9PB61;Qξ E9PB61;Qξ THO comp ALYREF
 0,682786 0,21492 E9PBC5;HC E9PBC5;HC Plasma kal KLKB1
 3,70332 0,002416 E9PCY7;G8 E9PCY7;G8 Heterogen HNRNPH1
 4,50725 0,000881 E9PEB5;Qξ E9PEB5;Qξ Far upstre FUBP1
 1,99563 0,023573 Q5TG12;Eξ Q5TG12;Eξ Protein-tyr PTPRK
 2,8926 0,006728 E9PGN7;PC E9PGN7;PC Plasma prc SERPING1
 1,55042 0,048112 E9PGT1;Qξ E9PGT1;Qξ Translin TSN
 4,06795 0,001521 E9PH05;F8 E9PH05;F8 Protein FAI FAM162A
 2,13681 0,018887 E9PH64;Qξ E9PH64;Qξ NADH deh' NDUFB9
 6,30885 0,000152 E9PHF7;Qξ E9PHF7;Qξ Methylcroi MCCC1
 3,3003 0,00394 E9PHK0;PC E9PHK0;PC Tetranecti CLEC3B
 6,78388 7,24E-05 E9PHY5;Oξ E9PHY5;Oξ Band 4.1-li EPB41L2
 1,83861 0,030013 P29692;E9 P29692;E9 Elongation EEF1D
 4,0524 0,001547 E9PLK3;P5 E9PLK3;P5 Puromycin NPEPPS
 5,13369 0,000472 E9PPW7;O E9PPW7;O NADH deh' NDUFS8
 2,6164 0,009519 F2Z3J9;Q1 F2Z3J9;Q1 Prostaglan PTGR1
 5,19737 0,000451 F5GYQ1;Pξ F5GYQ1;Pξ V-type pro ATP6V0D1
 9,87597 0 F5GZS6;J3I F5GZS6;J3I 4F2 cell-su SLC3A2
 2,62839 0,009364 F5H157;Q1 F5H157;Q1 Ras-relatec RAB35
 5,34155 0,000431 F5H365;Q1 F5H365;Q1 Protein tra SEC23A
 5,03552 0,000516 P55209;F8 P55209;F8 Nucleosom NAP1L1
 4,74974 0,000668 H0Y8R1;F5 H0Y8R1;F5 G-rich seqt GRSF1
 16,3597 0 F5H5P2;P1 F5H5P2;P1 2-oxoisova BCKDHA
 6,31074 0,000152 F5H6E2;OC F5H6E2;OC Unconvent MYO1C
 0,132781 0,683404 F6TLX2;Q9 F6TLX2;Q9 Glyoxalase GLOD4
 4,537 0,000853 P27708;F8 P27708;F8 CAD protei CAD
 6,75662 7,19E-05 F8VQR7;Q F8VQR7;Q Cysteine ai CSRP2
 1,21704 0,083207 F8VV04;Q1 F8VV04;Q1 Squamous SART3
 2,87867 0,006857 F8VV56;F8 F8VV56;F8 Tetraspani CD63
 3,16847 0,004645 F8VVA7;Pξ F8VVA7;Pξ Coatomer COPZ1
 11,5287 0 F8W1A4;P F8W1A4;P Adenylate AK2
 0,876243 0,150876 F8W914;Q F8W914;Q Reticulon;F RTN4
 0,853965 0,157835 J3QRU4;F8 J3QRU4;F8 Vesicle-ass VAMP2;VAMP3
 1,32313 0,069843 F8WCF6;Pξ F8WCF6;Pξ Actin-relat ARPC4-TTLL3;ARPC4
 3,21814 0,004386 G3V0E4;Oξ G3V0E4;Oξ Mitochond PMPCB
 13,0815 0 G3V0I5;P4 G3V0I5;P4 NADH deh' NDUFV1
 4,329 0,001123 G3V1D3;G G3V1D3;G Dipeptidyl DPP3
 1,94538 0,025521 G3V1A6;Pξ G3V1A6;Pξ Gasdermin GSDMD
 2,40254 0,012909 G3V1C3;Qξ G3V1C3;Qξ Apoptosis i API5
 5,25117 0,000444 G3V2S6;Qξ G3V2S6;Qξ V-type pro ATP6V1D
 3,44346 0,003378 G3V4U0;Q G3V4U0;Q Fibulin-5 FBLN5
 8,16235 0 P07942;G3 P07942;G3 Laminin su LAMB1
 4,07258 0,001502 G5E9Q6;Pξ G5E9Q6;Pξ Profilin;Prc PFN2
 0,604196 0,250275 O00429-6; O00429-6; Dynamin-1 DNM1L
 5,5765 0,000333 H0UI06;P1 H0UI06;P1 Cytochrom COX7A2
 5,20576 0,000453 P0DP25;PC P0DP25;PC CALM2;CALM1;CALM3
 3,73268 0,002367 H0Y8B3;Qξ H0Y8B3;Qξ Adenosylh AHCYL2

5,36309 0,000426 H0YF29;Q€ H0YF29;Q€ UPF0598 p C8orf82
1,95043 0,025285 H0YFA4;P5 H0YFA4;P5 Cysteine-ri CRIP2
1,15855 0,09172 H0YHA7;J3 H0YHA7;J3 60S riboso RPL18
1,54079 0,048783 Q9H8H3;H Q9H8H3;H Methyltrar METTL7A
0,25537 0,513497 H0YKD8;P4 H0YKD8;P4 60S riboso RPL28
3,26695 0,004123 H0YMZ1;H H0YMZ1;H Proteasom PSMA4
4,41253 0,001028 H3BNC9;P€ H3BNC9;P€ 40S riboso RPS17
5,91451 0,000243 H3BNT2;O` H3BNT2;O` Ubiquinon€ COQ9
1,70512 0,03718 H3BS10;H€ H3BS10;H€ Beta-hexo€ HEXA
3,19831 0,004502 H3BPE1;H€ H3BPE1;H€ Microtubu MACF1
3,32051 0,003825 H3BPK3;Q: H3BPK3;Q: Hydroxyac` HAGH
7,67001 1,95E-05 H3BQZ7;Q H3BQZ7;Q Heterogen HNRNPUL2-BSCL2;HNRNPUL2
13,529 0 H3BRN4;P€ H3BRN4;P€ 4-aminobu ABAT
6,03747 0,000193 H3BRY5;P€ H3BRY5;P€ Sulfotransf SULT1A1;SULT1A2;SULT1A4;SULT1A3
4,59354 0,000808 H3BSM0;Q H3BSM0;Q Acyl-coenz ACSM3
3,66239 0,002564 H3BUX2;O H3BUX2;O Cytochrom CYB5B
3,02561 0,005602 H7C003;Q: H7C003;Q: Protein ph` PPP1R7
19,3498 0 H7C1U8;Q H7C1U8;Q Apolipoprc APOO
1,59883 0,044385 H7C4H2;Q: H7C4H2;Q: Signal reco SRPRB
2,95571 0,006115 H7C5S0;O€ H7C5S0;O€ Actin-like € ACTL6A
3,07249 0,005309 I3L0N3;P4€ I3L0N3;P4€ Vesicle-fus NSF
0,497513 0,308952 I3L297 I3L297 CRK
4,87602 0,000588 J3KND8;P0 J3KND8;P0 Phenylalan PAH
6,39551 0,000132 J3KNQ4;Q€ J3KNQ4;Q€ Alpha-parv PARVA
10,0039 0 J3KPX7;Q9 J3KPX7;Q9 Prohibitin- PHB2
1,00363 0,1207 J3KQ32;Q€ J3KQ32;Q€ Obg-like A` OLA1
5,74919 0,00028 J3KQJ1;Q8 J3KQJ1;Q8 Sulfatase-r SUMF2
2,74024 0,00812 J3KR97;Q9 J3KR97;Q9 Tubulin-sp` TBCD
1,96766 0,024627 J3KTA4;P1` J3KTA4;P1` Probable A DDX5
0,771819 0,182933 J3KTF8;J3C J3KTF8;J3C Rho GDP-d ARHGDIA
2,93916 0,006259 J3KTL2;Q0` J3KTL2;Q0` Serine/argi` SRSF1
1,29214 0,073321 J3QLE5;P1· J3QLE5;P1· Small nucl€ SNRPN;SNRPB
4,07797 0,001496 J3QRS3;P1 J3QRS3;P1 Myosin re€ MYL12A;MYL12B
2,33343 0,014121 K7EIK7;O9` K7EIK7;O9` Echinoderr EML2
0,212791 0,564982 K7EJL1;Q9| K7EJL1;Q9| AP-1 comp AP1M1
0,505422 0,304058 K7EK07;P8 K7EK07;P8 Histone H3 H3F3B;H3F3A;HIST3H3;H3F3C
6,74537 7,14E-05 K7EQ77;K7 K7EQ77;K7 NADH deh` NDUFA11
3,35466 0,003674 K7EKI8;O6| K7EKI8;O6| Periplakin PPL
4,16848 0,00134 K7ELL7;P1· K7ELL7;P1· Glucosidas PRKCSH
4,52557 0,000878 K7EPR0;Q€ K7EPR0;Q€ Transmem TMEM205
12,5729 0 K7N7A8;P2 K7N7A8;P2 Aquaporin AQP1
3,19203 0,004522 M0QWZ7;€ M0QWZ7;€ Serine--tr` SARS2
3,40699 0,003502 M0QXB4;C M0QXB4;C Coatomer` COPE
1,43532 0,057759 O95571;M O95571;M Persulfide` ETHE1
2,68213 0,008771 M0R299;N M0R299;N rRNA 2-O-r FBL
4,03869 0,001571 P62280;M€ P62280;M€ 40S riboso RPS11
1,19321 0,086592 M0R0F0;P· M0R0F0;P· 40S riboso RPS5
4,36026 0,001081 M0R221;N M0R221;N U1 small n` SNRPA
0,511018 0,30081 O00116;AC O00116;AC Alkylidihyd€ AGPS
6,088 0,000182 O00151 O00151 PDZ and LI PDLIM1

0,713039 0,203697 O00231;O O00231;O 26S protea PSMD11
0,062331 0,82355 O00232;O O00232;O 26S protea PSMD12
1,53551 0,049067 O00264;O O00264;O Membrane PGRMC1
19,7371 0 O00299 O00299 Chloride in CLIC1
4,09062 0,001472 O00303;H O00303;H Eukaryotic EIF3F
12,3639 0 O00330-3; O00330-3; Pyruvate d PDHX
0,960069 0,129966 O00410;O O00410;O Importin-5 IPO5
3,79152 0,002181 O00468-6; O00468-6; Agrin;Agrir AGRN
3,85188 0,002 O00483 O00483 Cytochrom NDUFA4
2,53648 0,010726 O00487;C O00487 26S protea PSMD14
1,96825 0,02461 O00567;A O00567;A Nucleolar r NOP56
2,68576 0,008738 O00592-2; O00592-2; Podocalyxin PODXL
3,81419 0,002094 O00629;H O00629 Importin s1 KPNA4
9,03032 0 O00748;O O00748;O Cocaine est CES2
2,22589 0,016474 O00764;F2 O00764;F2 Pyridoxal k PDXK
3,63734 0,002627 O14548;E5 O14548;E5 Cytochrom COX7A2L
7,88545 2,03E-05 O14638;F8 O14638;F8 Ectonucleoside ENPP3
3,05529 0,00544 O14732-2; O14732-2; Inositol m1 IMPA2
14,5083 0 O14745;J3 O14745;J3 Na(+)/H(+) SLC9A3R1
1,75178 0,034509 O14818;O O14818;O Proteasome PSMA7
4,86308 0,00059 O14880;Q O14880;Q Microsomal MGST3
20,3023 0 O14949;D O14949 Cytochrom UQCRCQ
3,31705 0,003833 O14957 O14957 Cytochrom UQCR11
11,3783 0 O14975;O O14975;O Very long-c SLC27A2
0,475006 0,322146 O14979-3; O14979-3; Heterogeneous HNRNPDL
4,24275 0,00123 O14980;C O14980 Exportin-1 XPO1
5,43883 0,000411 O15020;O O15020;O Spectrin b1 SPTBN2
2,23899 0,01617 O15067;J3 O15067 Phosphoribosyl PFAS
5,16771 0,000459 O15143;C O15143;C Actin-related ARPC1B
1,05049 0,111176 O15144;H O15144 Actin-related ARPC2
2,03839 0,022157 O15145;C O15145;C Actin-related ARPC3
5,17454 0,000448 O15230;O O15230 Laminin subunit LAMA5
0,350846 0,41685 Q5T8U5;O Q5T8U5;O Surfeit locus SURF4
4,8943 0,00058 O15511;B1 O15511;B1 Actin-related ARPC5
1,31669 0,07056 O15540;O O15540;O Fatty acid-binding FABP7
7,56948 2,82E-05 O43237;O O43237;O Cytoplasmic DYNC1L12
2,91479 0,006497 O43242;O O43242;O 26S protea PSMD3
4,56498 0,000836 O43294;O O43294;O Transforming TGFB111
6,01684 0,000206 O43301;A O43301;A Heat shock HSPA12A
0,810449 0,170371 O43390;O O43390;O Heterogeneous HNRNP
3,0913 0,005129 O43396;K7 O43396;K7 Thioredoxin TXNL1
6,36355 0,000138 O43488;H O43488;H Aflatoxin B AKR7A2
7,79007 2E-05 O43570-2; O43570-2; Carbonic anhydrase CA12
5,2897 0,000447 O43598;H O43598;H 2-deoxyribose DNPH1
4,1041 0,00145 O43615;M O43615;M Mitochondrial TIMM44
3,19341 0,004518 O43681;A O43681;A ATPase subunit ASNA1
10,5359 0 O43707;O O43707;O Alpha-actinin ACTN4
1,10104 0,101427 O43747;O O43747;O AP-1 component AP1G1
1,32662 0,069475 O43772;C O43772;C Mitochondrial SLC25A20
1,85152 0,029389 O43776;K7 O43776 Asparagine NARS

0,269594 0,497057 O43813;E9 O43813;E9 LanC-like p LANCL1
5,62535 0,000315 O43865-2; O43865-2; Putative ac AHCYL1
4,9771 0,000534 O43895 O43895 Xaa-Pro an XPNPEP2
5,7097 0,000292 O60343-2; O60343-2; TBC1 dom TBC1D4
1,90986 0,026818 O60493-4; O60493-4; Sorting ne SNX3
7,61511 1,91E-05 O60494;H O60494 Cubilin CUBN
2,16214 0,018082 O60506-3; O60506-3; Heterogen SYNCRIP
2,57038 0,010147 O60568;H O60568 Procollage PLOD3
14,4781 0 O60656;O O60656;O UDP-glucu UGT1A9;UGT1A8;UGT1A5;UGT1A3;UGT1A4;
1,70654 0,037119 O60664-4; O60664-4; Perilipin-3 PLIN3
1,25782 0,077832 O60684;Q O60684;Q Importin s KPNA6
8,13224 0 O60701-2; O60701-2; UDP-glucu UGDH
2,41415 0,012778 O60749;O O60749;O Sorting ne SNX2
2,83774 0,007169 O60763;O O60763;O General ve USO1
6,75964 0,000072 O60784;O O60784;O Target of M TOM1
1,48069 0,053603 U3KQK0;Q U3KQK0;Q Histone H2 HIST1H2BN;HIST1H2BM;HIST1H2BH;HIST2H;
0,132694 0,683112 O60884;A O60884 Dnal homc DNAJA2
1,16948 0,089982 O60888-3; O60888-3; Protein Cu CUTA
11,5823 0 O75083;O O75083;O WD repeat WDR1
1,01268 0,119064 O75131;A O75131;A Copine-3 CPNE3
0,721417 0,200477 O75146;O O75146;O Huntingtin HIP1R
11,1509 0 O75251;F5 O75251;F5 NADH deh NDUFS7
7,66993 1,95E-05 O75306;O O75306;O NADH deh NDUFS2
11,9503 0 O75309-2; O75309-2; Cadherin-1 CDH16
15,9672 0 O75323;O O75323;O Protein Ni GBAS
3,02149 0,005623 O75340-2; O75340-2; Programm PDCD6
3,26996 0,004106 O75348;F2 O75348 V-type pro ATP6V1G1
2,84161 0,007179 O75367-2; O75367-2; Core histor H2AFY
3,73958 0,002335 O75369-8; O75369-8; Filamin-B FLNB
0,724062 0,199878 O75396;A O75396 Vesicle-tra SEC22B
3,21432 0,004394 O75431;O O75431;O Metaxin-2 MTX2
0,980161 0,125392 O75436;S4 O75436 Vacuolar p VPS26A
6,4468 0,00011 O75489;E9 O75489 NADH deh NDUFS3
2,4585 0,011963 O75531 O75531 Barrier-to BANF1
1,1095 0,100051 O75533;H O75533 Splicing fac SF3B1
2,12654 0,019197 O75608-2; O75608-2; Acyl-protei LYPLA1
8,12853 0 O75643;A O75643;A U5 small n SNRNP200
10,8432 0 O75746;O O75746;O Calcium-bi SLC25A12
7,26629 3,53E-05 O75795;A O75795;A UDP-glucu UGT2B17;UGT2B15
8,17235 0 O75874;C9 O75874 Isocitrate c IDH1
5,97747 0,000225 O75880;J3 O75880;J3 Protein SC SCO1
3,41475 0,003497 O75891;O O75891;O Cytosolic 1 ALDH1L1
4,70843 0,00069 O75915;F8 O75915;F8 PRA1 famil ARL6IP5
5,39148 0,000427 O75923-1E O75923-1E Dysferlin DYSF
11,1944 0 O75936;E9 O75936 Gamma-bu BBOX1
8,10092 1,04E-05 O75947;O O75947;O ATP synth ATP5H
5,16447 0,000458 O75955;O O75955;O Flotillin-1 FLOT1
9,6319 0 O75964;E9 O75964;E9 ATP synth ATP5L
1,48784 0,053022 O76038;Q O76038 Secretago SCGN
9,75022 0 O94760;B4 O94760;B4 N(G),N(G)- DDAH1

2,333	0,01413	O94826	O94826	Mitochond	TOMM70A
2,25987	0,01568	O94905;E5	O94905;E5	Erlin-2	ERLIN2
3,91842	0,001807	O94919	O94919	Endonucle	ENDOD1
5,40348	0,000428	O94925;B7	O94925	Glutamina:	GLS
6,52166	0,000107	O95154	O95154	Aflatoxin B	AKR7A3
7,27456	3,55E-05	O95168;O!	O95168;O!	NADH deh	NDUFB4
3,84196	0,002017	O95169-3;	O95169-3;	NADH deh	NDUFB8
4,06478	0,001527	O95197-3;	O95197-3;	Reticulon-	RTN3
8,55465	0	O95202;O!	O95202	LETM1 anc	LETM1
1,72383	0,036079	O95336;M	O95336;M	6-phospho	PGLS
14,2745	0	Q5R3B4;O!	Q5R3B4;O!	Mitochond	MPC2
0,554217	0,276414	O95573;O!	O95573	Long-chain	ACSL3
0,924038	0,138157	O95822;O!	O95822;O!	Malonyl-C	MLYCD
14,3373	0	O95831;O!	O95831;O!	Apoptosis-	AIFM1
2,37784	0,013349	O95865;AC	O95865;AC	N(G),N(G)-	DDAH2
9,81943	0	O95954;O!	O95954;O!	Formimido	FTCD
9,72689	0	O96008;O!	O96008;O!	Mitochond	TOMM40
3,92385	0,001795	O96009;M	O96009;M	Napsin-A	NAPSA
16,3985	0	P00167;P0	P00167;P0	Cytochrom	CYB5A
7,53933	2,82E-05	P00325;V9	P00325;V9	Alcohol de	ADH1B;HEL-S-117;ADH1A
14,5954	0	P00338;P0	P00338;P0	L-lactate d	LDHA
3,7067	0,002409	P00352;Q!	P00352	Retinal de	ALDH1A1
12,63	0	P00367;P0	P00367;P0	Glutamate	GLUD1;GLUD2
2,35187	0,013819	P00387-2;I	P00387-2;I	NADH-cytc	CYB5R3
0,545442	0,281404	P00390-2;I	P00390-2;I	Glutathion	GSR
2,96114	0,006068	P00395;DI!	P00395	Cytochrom	MT-CO1
8,21127	0	P00403	P00403	Cytochrom	MT-CO2
9,44732	0	P00414	P00414	Cytochrom	MT-CO3
5,88232	0,000249	P00441;H7	P00441;H7	Superoxide	SOD1
0,648131	0,229948	P00450;E9	P00450;E9	Ceruloplas	CP
10,0851	0	P00488;Q!	P00488	Coagulatio	F13A1
7,29159	3,56E-05	P00491;G3	P00491;G3	Purine nuc	PNP
12,5013	0	P00492	P00492	Hypoxanth	HPRT1
16,5801	0	P00505;P0	P00505;P0	Aspartate	GOT2
8,90712	0	P00558;P0	P00558;P0	Phosphogl	PGK1
2,13525	0,018909	Q5T9B7;PC	Q5T9B7;PC	Adenylate	AK1
2,72552	0,008298	P00734;E9	P00734;E9	Prothromb	F2
0,656203	0,226429	P00738;J3!	P00738;J3!	Haptoglobi	HP
3,87275	0,001957	P00740;P0	P00740;P0	Coagulatio	F9
2,41951	0,01268	P00747;Q!	P00747	Plasminog	PLG
5,80343	0,000268	P00846;DI!	P00846	ATP synth	MT-ATP6
1,81126	0,031393	P00915;E5	P00915;E5	Carbonic a	CA1
10,547	0	P00918;E5	P00918	Carbonic a	CA2
15,2613	0	P00966;Q!	P00966;Q!	Argininosu	ASS1
1,47064	0,054549	P01008;Q!	P01008	Antithromb	SERPINC1
2,6158	0,009521	P01009;A0	P01009;A0	Alpha-1-an	SERPINA1
0,526662	0,291748	P01011;G3	P01011;G3	Alpha-1-an	SERPINA3
3,60561	0,002722	P01019	P01019	Angiotensi	AGT
0,314995	0,450292	P01023;F8	P01023	Alpha-2-m	A2M
1,58743	0,045206	P01024;M!	P01024	Compleme	C3

2,10774	0,019854	P01031;DI	P01031	Compleme C5
2,53002	0,010817	P01042;P0	P01042;P0	Kininogen- KNG1
11,2735	0	P01833;DI	P01833	Polymeric iPIGR
1,94213	0,025603	P01834;DI	P01834	Ig kappa cł IGKC
0,620284	0,242592	P02042;E9	P02042;E9	Hemoglobi HBD
1,37383	0,064105	P02452	P02452	Collagen al COL1A1
7,31501	3,59E-05	P02511;E9	P02511;E9	Alpha-crysi CRYAB
4,65755	0,000759	P02545;P0	P02545;P0	Prelamin-A LMNA
2,59695	0,009774	P02545-2;(P02545-2;(Prelamin-A LMNA
0,525172	0,292459	P02549-2;I	P02549-2;I	Spectrin al SPTA1
7,29742	3,57E-05	P02647;F8	P02647;F8	Apolipoprc APOA1
0,491544	0,312415	P02649;HC	P02649;HC	Apolipoprc APOE
3,76975	0,00225	V9GYM3;P	V9GYM3;P	Apolipoprc APOA2
0,022963	0,925474	P02671-2;I	P02671-2;I	Fibrinogen FGA
0,072848	0,8	P02675;D6	P02675;D6	Fibrinogen FGB
0,444961	0,342157	P02679-2;I	P02679-2;I	Fibrinogen FGG
1,59414	0,044756	P02730;P0	P02730;P0	Band 3 ani SLC4A1
3,97265	0,00167	P02743	P02743	Serum amy APCS
0,769591	0,183509	P02747	P02747	Compleme C1QC
2,87522	0,006871	P02748	P02748	Compleme C9
1,59111	0,044959	P02749;J3(P02749;J3(Beta-2-glyc APOH
4,27151	0,001181	P02751-15	P02751-15	Fibronectir FN1
6,9572	6,63E-05	Q5VY30;P(Q5VY30;P(Retinol-bin RBP4
2,84144	0,007172	P02760;S4	P02760;S4	Protein AN AMBP
0,775231	0,181951	P02763	P02763	Alpha-1-ac ORM1
10,2972	0	P02765;C9	P02765;C9	Alpha-2-HS AHSG
7,62183	1,93E-05	P02766;A0	P02766;A0	Transthyre TTR
7,47903	2,77E-05	P02787;H7	P02787	Serotransfi TF
3,2705	0,00411	P02790;Q5	P02790;Q5	Hemopexir HPX
0,644511	0,231444	P02792	P02792	Ferritin ligł FTL
3,92751	0,001783	P02794;G3	P02794;G3	Ferritin heł FTH1
0,966514	0,12854	P03989	P03989	HLA class I HLA-B
0,597869	0,253165	P04003;A6	P04003	C4b-bindin C4BPA
3,57028	0,002858	P04004;F5	P04004	Vitronectir VTN
9,89608	0	P04040;DI	P04040	Catalase CAT
3,37958	0,003578	P04075;J3I	P04075;J3I	Fructose-b ALDOA
0,217643	0,559103	P04080;A0	P04080;A0	Cystatin-B CSTB
7,22717	4,38E-05	P04083;Q5	P04083	Annexin A1 ANXA1
2,34778	0,01389	P04114;A8	P04114	Apolipoprc APOB
3,48881	0,003181	P04179;P0	P04179;P0	Superoxide SOD2
5,39146	0,000426	P04181;P0	P04181;P0	Ornithine ε OAT
8,16963	0	P04196	P04196	Histidine-ri HRG
3,7457	0,002331	P04217;P0	P04217;P0	Alpha-1B-g A1BG
4,87982	0,000589	P04229;P7	P04229	HLA class II HLA-DRB1
10,5894	0	P04275;P0	P04275	von Willeb VWF
4,19373	0,001282	P04350;MI	P04350	Tubulin be TUBB4A
9,66336	0	P04406;E7	P04406;E7	Glyceralde GAPDH
4,00065	0,001621	P04424-2;I	P04424-2;I	Argininosu ASL
12,0111	0	P04792;C9	P04792	Heat shock HSPB1
9,74114	0	P04843;B7	P04843;B7	Dolichyl-di RPN1

5,13852 0,000473 P04844;P0 P04844;P0 Dolichyl-di RPN2
6,58733 8,56E-05 P04899-4;I P04899-4;I Guanine n GNAI2
0,202446 0,578293 Q93077;Q Q93077;Q Histone H2 HIST1H2AC;HIST3H2A;HIST1H2AB
24,6233 0 P05023-4;I P05023-4;I Sodium/pc ATP1A1
15,5702 0 P05026;P0 P05026;P0 Sodium/pc ATP1B1
9,91592 0 P05091;P0 P05091;P0 Aldehyde c ALDH2
2,49465 0,011338 P05109 P05109 Protein S1(S100A8
15,9764 0 P05141 P05141 ADP/ATP t SLC25A5
6,93063 6,58E-05 P05154;G3 P05154 Plasma ser SERPINA5
0,828073 0,16531 P05164-2;I P05164-2;I Myeloperc MPO
9,93225 0 P05165-2;I P05165-2;I Propionyl- PCCA
13,8449 0 P05166;E9 P05166;E9 Propionyl- PCCB
4,00247 0,001623 P05198;HC P05198;HC Eukaryotic EIF2S1
6,74831 7,16E-05 P05386;HC P05386 60S acidic RPLP1
1,79852 0,03194 P05388;F8 P05388;F8 60S acidic RPLP0;RPLPOP6
0,93487 0,135555 S4R371;P0 S4R371;P0 Fatty acid- FABP3
13,5714 0 P05556;P0 P05556;P0 Integrin be ITGB1
14,0924 0 P05783;F8 P05783;F8 Keratin, ty KRT18
13,146 0 P05937-2;I P05937-2;I Calbindin CALB1
1,13626 0,095237 P05976-2;I P05976-2;I Myosin lig MYL1;MYL3
14,5567 0 P06396-2;I P06396-2;I Gelsolin GSN
16,8798 0 P06576;HC P06576;HC ATP synth ATP5B
2,98731 0,005897 P06703;R4 P06703;R4 Protein S1(S100A6
2,29915 0,014821 P06727 P06727 Apolipoprc APOA4
15,6322 0 P06733;A0 P06733;A0 Alpha-enol ENO1
9,09616 0 P06737-2;I P06737-2;I Glycogen P PYGL
4,96611 0,000538 P06744;A0 P06744;A0 Glucose-6- GPI
4,75296 0,000669 P06748-2;I P06748-2;I Nucleopho NPM1
1,79463 0,032086 P06756-3;I P06756-3;I Integrin al ITGAV
1,64591 0,040927 Q16778;P Q16778;P Histone H2 HIST2H2BE;HIST1H2BB;HIST1H2BO;HIST1H2I
3,51291 0,003091 P07099;B1 P07099 Epoxide hy EPHX1
6,83462 6,5E-05 P07195;A0 P07195;A0 L-lactate d LDHB
19,6715 0 P07237;H7 P07237;H7 Protein dis P4HB
11,4052 0 P07355;P0 P07355;P0 Annexin A ANXA2
2,24126 0,016131 P07358;F5 P07358;F5 Compleme C8B
13,7275 0 P07384;E9 P07384 Calpain-1 c CAPN1
14,7003 0 Q5JP53;P0 Q5JP53;P0 Tubulin be TUBB
7,36623 2,73E-05 P07585;P0 P07585;P0 Decorin DCN
2,69911 0,00857 Q5URX0;P Q5URX0;P Beta-hexo HEXB
8,67968 0 P07737;K7 P07737;K7 Profilin-1 PFN1
1,78676 0,032441 P07741;H3 P07741;H3 Adenine p APRT
1,07659 0,105911 P07814;V9 P07814 Bifunction EPRS
11,1437 0 P07858;R4 P07858 Cathepsin I CTSB
3,75297 0,002295 P07900;P0 P07900;P0 Heat shock HSP90AA1
16,7522 0 X6RBG4;P X6RBG4;P Uromoduli UMOD
8,26936 0 P07919;A0 P07919;A0 Cytochrom UQCRH;UQCRHL
14,5057 0 P07954;P0 P07954;P0 Fumarate I FH
5,04993 0,000507 P07996;P0 P07996;P0 Thrombos THBS1
6,74991 7,17E-05 P08133;P0 P08133;P0 Annexin A ANXA6
3,97168 0,001666 P08183;P0 P08183;P0 Multidrug ABCB1

3,5672	0,002859	P08236;P0	P08236;P0	Beta-glucu	GUSB
3,4424	0,003387	P08238;Q5	P08238	Heat shock	HSP90AB1
1,78784	0,032422	P08246	P08246	Neutrophil	ELANE
14,9527	0	P08263;Q1	P08263	Glutathion	GSTA1
11,2442	0	P08473;C9	P08473	Neprilysin	MME
16,5763	0	P08559;P0	P08559;P0	Pyruvate d	PDHA1
2,26197	0,01563	P08571;D6	P08571;D6	Monocyte	CD14
3,51042	0,003092	P08572;A0	P08572;A0	Collagen al	COL4A2
14,9678	0	P08574;D1	P08574	Cytochrom	CYC1
4,21815	0,001249	P08603;P0	P08603	Compleme	CFH
14,7321	0	P08670;B0	P08670;B0	Vimentin	VIM
2,80979	0,007443	P08754	P08754	Guanine n	GNAI3
12,1558	0	P08758;D6	P08758;D6	Annexin A5	ANXA5
9,05786	0	P09104;P0	P09104;P0	Gamma-er	ENO2
12,9717	0	P09110;C9	P09110;C9	3-ketoacyl	ACAA1
9,36962	0	P09210	P09210	Glutathion	GSTA2
4,39243	0,001055	P09211;A8	P09211;A8	Glutathion	GSTP1
7,57803	2,83E-05	P09327;B4	P09327;B4	Villin-1	VIL1
10,5859	0	P09382;F8	P09382	Galectin-1	LGALS1
5,16188	0,000457	P09417;P0	P09417;P0	Dihydropte	QDPR
0,231621	0,541574	P09429;Q5	P09429;Q5	High mobil	HMGB1
15,801	0	P09467;A0	P09467;A0	Fructose-1	FBP1
15,746	0	P09525;Q6	P09525;Q6	Annexin A4	ANXA4
4,88933	0,000585	P09543-2;I	P09543-2;I	2,3-cyclic-r	CNP
19,383	0	P09622;E9	P09622;E9	Dihydrolipi	DLD
9,31216	0	P09651-2;I	P09651-2;I	Heterogen	HNRNPA1
5,52711	0,000374	P09669	P09669	Cytochrom	COX6C
1,39029	0,062365	P09874	P09874	Poly [ADP-	PARP1
5,23281	0,000448	P09936;D6	P09936;D6	Ubiquitin c	UCHL1
6,51318	0,000106	P09960;P0	P09960;P0	Leukotrien	LTA4H
8,76056	0	P09972;A8	P09972;A8	Fructose-b	ALDOC
2,97449	0,005988	P0C0L4	P0C0L4	Compleme	C4A
4,23741	0,001234	P0DN76;Q1	P0DN76;Q1	Splicing fac	U2AF1;U2AF1L4
1,38616	0,062718	P0DOY2;P0	P0DOY2;P0		
1,68322	0,038491	P10155-3;I	P10155-3;I	60 kDa SS-	TROVE2
5,24539	0,000443	P10176;U3	P10176	Cytochrom	COX8A
5,64033	0,00031	P10253;I3	P10253	Lysosomal	GAA
6,3688	0,000139	P10301	P10301	Ras-relate	RRAS
3,04402	0,005486	P10412	P10412	Histone H1	HIST1H1E
18,2009	0	P10515;E9	P10515;E9	Dihydrolipi	DLAT
0,479799	0,319419	P10599;P1	P10599;P1	Thioredoxi	TXN
5,13405	0,000472	P10606	P10606	Cytochrom	COX5B
2,55338	0,010426	X6R5C5;X6	X6R5C5;X6	Carboxype	CTSA
0,649652	0,229446	P10644;P1	P10644;P1	cAMP-dep	PRKAR1A
6,56778	9,25E-05	P10768;X6	P10768;X6	S-formylgl	ESD
9,66203	0	P10809;E7	P10809	60 kDa he	HSPD1
2,80641	0,007483	P10909-4;I	P10909-4;I	Clusterin;	CLU
5,52027	0,000373	P11021	P11021	78 kDa glu	HSPA5
4,13816	0,001403	P11047;R4	P11047	Laminin su	LAMC1
1,89101	0,027603	P11142;E9	P11142;E9	Heat shock	HSPA8

1,58697	0,045205	P11172;E9	P11172;E9	Uridine 5-r UMPS
12,734	0	P11177-2;I	P11177-2;I	Pyruvate d PDHB
7,05228	5,06E-05	P11182;Q5	P11182;Q5	Lipoamide DBT
9,24018	0	P11216;HC	P11216	Glycogen ç PYGB
7,097	5,1E-05	P11233;H7	P11233;H7	Ras-relatec RALA
0,185531	0,601665	P11277;P1	P11277;P1	Spectrin bε SPTB
5,98566	0,000219	P11279;P1	P11279;P1	Lysosome- LAMP1
5,79331	0,000267	Q5T4U5;P:	Q5T4U5;P:	Medium-cl ACADM
9,03234	0	P11413;P1	P11413;P1	Glucose-6- G6PD
10,6131	0	P11498;A0	P11498;A0	Pyruvate c PC
1,19368	0,086539	P11717	P11717	Cation-indı IGF2R
3,73646	0,002346	P11766;HC	P11766;HC	Alcohol de ADH5
1,55258	0,047971	P12074;HC	P12074;HC	Cytochrom COX6A1
8,74361	0	P12110;P1	P12110;P1	Collagen al COL6A2
12,8567	0	P12111-2;I	P12111-2;I	Collagen al COL6A3
9,74667	0	P12111-4;I	P12111-4;I	Collagen al COL6A3
12,3582	0	P12235;V9	P12235;V9	ADP/ATP tı SLC25A4
14,4609	0	P12236	P12236	ADP/ATP tı SLC25A6
0,629188	0,238551	P12270;P1	P12270;P1	Nucleopro: TPR
16,031	0	P12277;HC	P12277;HC	Creatine ki CKB
2,61474	0,009539	P12429;D6	P12429;D6	Annexin A3 ANXA3
16,4021	0	P12532;P1	P12532;P1	Creatine ki CKMT1A;CKMT1B
9,81195	0	P12814-3;I	P12814-3;I	Alpha-actir ACTN1
6,87391	6,54E-05	P12955;K7	P12955;K7	Xaa-Pro diı PEPD
1,4203	0,059242	P12956;B1	P12956;B1	X-ray repai XRCC6
7,72628	1,97E-05	P13010;C9	P13010	X-ray repai XRCC5
7,89458	2,04E-05	P13073;H3	P13073;H3	Cytochrom COX4I1
3,74271	0,002338	P13284	P13284	Gamma-inı IFI30
11,0177	0	P13473-2;I	P13473-2;I	Lysosome- LAMP2
2,21338	0,016746	P13489;HC	P13489	Ribonuclea RNH1
8,96086	0	P13611-4;I	P13611-4;I	Versican cç VCAN
13,296	0	P13639;DI	P13639	Elongation EEF2
13,353	0	P13667;C9	P13667	Protein dis PDIA4
2,12162	0,019371	P13674-3;I	P13674-3;I	Prolyl 4-hy P4HA1
6,06648	0,000188	P13716;P1	P13716;P1	Delta-amir ALAD
0,377146	0,394043	P13746;P1	P13746;P1	HLA class I HLA-A
11,2083	0	P13796;P1	P13796	Plastin-2 LCP1
11,1662	0	P13804;HC	P13804;HC	Electron tr ETFA
6,17541	0,000178	P13807-2;I	P13807-2;I	Glycogen [GYS1
0,75577	0,188168	P13861-2;I	P13861-2;I	cAMP-depı PRKAR2A
0,593903	0,254956	P14174	P14174	Macrophagı MIF
11,0184	0	P14543;P1	P14543;P1	Nidogen-1 NID1
10,7798	0	P14550;Q5	P14550	Alcohol de AKR1A1
9,67902	0	P14618;P1	P14618;P1	Pyruvate k PKM
12,2327	0	P14625;Q5	P14625	Endoplasm HSP90B1
9,22676	0	P14854;A0	P14854	Cytochrom COX6B1
4,43629	0,000985	P14866;Mı	P14866;Mı	Heterogen HNRNPL
5,26338	0,000444	P14868;P1	P14868;P1	Aspartate- DARS
6,20179	0,000178	P14923;C9	P14923	Junction pl JUP
15,9193	0	P14927;B7	P14927;B7	Cytochrom UQCRB

1,55533 0,047753 P15121;E9 P15121;E9 Aldose red AKR1B1
14,1088 0 P15144;HC P15144 Aminopept ANPEP
3,85086 0,002002 P15153;B1 P15153;B1 Ras-related RAC2
15,3728 0 P15313;C9 P15313;C9 V-type pro ATP6V1B1
2,34535 0,013912 P15880;HC P15880;HC 40S riboso RPS2
4,13576 0,001399 P15924;P1 P15924;P1 Desmoplak DSP
4,4165 0,001031 P15954;D6 P15954;D6 Cytochrom COX7C
6,0675 0,000188 Q5TD07;P1 Q5TD07;P1 Ribosyldih NQO2
1,44701 0,056768 P16152;E9 P16152 Carbonyl re CBR1
0,367151 0,402595 P16157-14 P16157-14 Ankyrin-1 ANK1
13,227 0 P16219;E9 P16219;E9 Short-chain ACADS
7,26718 3,54E-05 P16278-3;I P16278-3;I Beta-galact GLB1
4,25874 0,001211 P16284-5;I P16284-5;I Platelet en PECAM1
0,194648 0,588777 P16403;P2 P16403 Histone H1 HIST1H1C
5,21944 0,000447 P16435;HC P16435;HC NADPH--cy POR
17,2692 0 P16444;H3 P16444 Dipeptidas DPEP1
11,1616 0 P16615;P1 P16615;P1 Sarcoplasn ATP2A2
14,7568 0 P16662;E9 P16662;E9 UDP-glucosyl UGT2B7
21,376 0 P16885;A0 P16885;A0 1-phosphatidyl PLCG2
5,01139 0,000519 P16930;P1 P16930;P1 Fumarylacetyl FAH
6,82701 6,49E-05 P17066;P4 P17066 Heat shock HSPA6
8,41291 0 P17174;P1 P17174;P1 Aspartate aminotransferase GOT1
0,994098 0,122467 P17252;J3I P17252;J3I Protein kinase PRKCA
19,0676 0 P17540;D6 P17540 Creatine kinase CKMT2
4,30833 0,001139 P17655;P1 P17655;P1 Calpain-2 class II CAPN2
3,25798 0,004161 P17661;D1I P17661 Desmin DES
8,69319 0 P17858;P1 P17858;P1 ATP-dependent PFKFB3
4,1088 0,001442 P17900;HC P17900 Ganglioside GM2A
7,66346 1,94E-05 P17931;G3 P17931;G3 Galectin-3 LGALS3
0,273479 0,493001 P17987;E7 P17987;E7 T-complex TCP1
0,479406 0,319455 P18124;A8 P18124;A8 60S ribosomal RPL7
3,66327 0,002548 P18206-2;I P18206-2;I Vinculin VCL
0,524049 0,292966 P18669;P1 P18669 Phosphoglycosyl PGAM1
4,33875 0,001109 P18754;P1 P18754;P1 Regulator of GSK-3B RGS1
6,29801 0,000159 P19338;H7 P19338;H7 Nucleolin NCL
10,9275 0 P19367;P1 P19367;P1 Hexokinase HK1
9,83772 0 P19440;E7 P19440;E7 Gamma-glutamyl GGT1;GGT2;GGT3P
9,04422 0 P19801;P1 P19801;P1 Amiloride receptor AOC1
1,31601 0,070613 P19823;Q5 P19823;Q5 Inter-alpha1(I) collagen ITIH2
7,76314 1,98E-05 P19827;P1 P19827;P1 Inter-alpha1(I) collagen ITIH1
13,1859 0 P19971;P1 P19971;P1 Thymidine kinase TYMP
5,00213 0,000518 P20073-2;I P20073-2;I Annexin A7 ANXA7
1,4349 0,057752 P20340-2;I P20340-2;I Ras-related RAB6A
1,57146 0,046344 P20618 P20618 Proteasome PSMB1
0,471246 0,324404 P20645;F5 P20645 Cation-dependent M6PR
10,9813 0 P20674;H3 P20674;H3 Cytochrom COX5A
12,9395 0 P20700;E9 P20700;E9 Lamin-B1 LMNB1
6,12813 0,000176 P20774;Q5 P20774;Q5 Mimecan OGN
1,74163 0,035069 P20933;HC P20933;HC N(4)-(beta)-mannosidase AGA
3,62101 0,002671 P21266;A0 P21266;A0 Glutathione GSTM3

12,3197	0	P21281;HC P21281	V-type pro ATP6V1B2
0,365508	0,40377	P21283;E7 P21283	V-type pro ATP6V1C1
7,01084	5,86E-05	P21291;E9 P21291;E9	Cysteine ar CSRP1
8,72955	0	P21333-2;I P21333-2;I	Filamin-A FLNA
7,18257	4,33E-05	P21397;P2 P21397;P2	Amine oxid MAOA
7,81033	0,00002	P21399 P21399	Cytoplasm ACO1
14,6132	0	P21695;P2 P21695;P2	Glycerol-3- GPD1
14,0577	0	P21796;C9 P21796;C9	Voltage-de VDAC1
8,68488	0	P21810;C9 P21810	Biglycan BGN
10,029	0	P21912;A0 P21912;A0	Succinate c SDHB
9,35193	0	P21953;P2 P21953	2-oxoisova BCKDHB
2,89648	0,006711	P21964-2;I P21964-2;I	Catechol O COMT
11,9396	0	P21980;P2 P21980;P2	Protein-glu TGM2
4,81062	0,000608	P22033 P22033	Methylmal MUT
3,86647	0,001964	P22059;HC P22059	Oxysterol-I OSBP
1,38365	0,062971	P22102;F8 P22102;F8	Trifunction GART
6,27462	0,000157	P22234;E9 P22234;E9	Multifunct PAICS
4,30139	0,001143	P22307-8;I P22307-8;I	Non-specif SCP2
5,12973	0,000471	P22314-2;I P22314-2;I	Ubiquitin-I UBA1
4,18065	0,001298	Q32Q12;P: Q32Q12;P:	Nucleoside NME1-NME2;NME2;NME1
5,05001	0,000508	P22626;P2 P22626;P2	Heterogen HNRNPA2B1
18,0767	0	P22695;H3 P22695;H3	Cytochrom UQCRC2
5,11956	0,00047	P23142;P2 P23142;P2	Fibulin-1 FBLN1
2,35198	0,013828	P23229-4;I P23229-4;I	Integrin al β ITGA6
2,40992	0,012839	P23246;P2 P23246;P2	Splicing fac SFPQ
2,84668	0,007154	P23284;DI: P23284	Peptidyl-pr PPIB
8,1862	0	P23381;P2 P23381;P2	Tryptophan WARS
10,6495	0	P23396;P2 P23396;P2	40S riboso RPS3
8,75162	0	P23434;A0 P23434;A0	Glycine cle GCSH
2,31368	0,014488	P23526;P2 P23526;P2	Adenosylh α AHCY
4,56167	0,00084	P23528;E9 P23528;E9	Cofilin-1 CFL1
2,26199	0,015644	P23786;A0 P23786;A0	Carnitine C CPT2
5,94836	0,000231	P24310;K7 P24310	Cytochrom COX7A1
2,3925	0,013057	P24534;C9 P24534	Elongation EEF1B2
10,5585	0	P24539;Q5 P24539;Q5	ATP synth α ATP5F1
2,51377	0,011017	P24666;G5 P24666;G5	Low molec ACP1
14,0967	0	P24752;HC P24752	Acetyl-CoA ACAT1
6,04954	0,000194	P24821-4;I P24821-4;I	Tenascin TNC
5,62756	0,000315	P24844;P2 P24844;P2	Myosin re ξ MYL9
2,75996	0,007903	P25325-2;I P25325-2;I	3-mercapt α MPST
0,416102	0,362918	P25391 P25391	Laminin su LAMA1
0,296	0,468928	P25398 P25398	40S riboso RPS12
15,9253	0	P25705;P2 P25705;P2	ATP synth α ATP5A1
1,79482	0,032106	P25786;P2 P25786;P2	Proteasom PSMA1
3,90753	0,001844	P25788-2;I P25788-2;I	Proteasom PSMA3
6,9856	6,67E-05	P26006;P2 P26006;P2	Integrin al β ITGA3
6,94714	6,6E-05	P26038;E9 P26038	Moesin MSN
4,73761	0,000667	P26368;P2 P26368;P2	Splicing fac U2AF2
4,91491	0,000558	P26373;J3(P26373;J3(60S riboso RPL13
9,79544	0	P26599;P2 P26599;P2	Polypyrimi PTBP1

1,02188 0,117115 P26639;P2 P26639;P2 Threonine- TARS
4,91696 0,000553 P26640;A0 P26640;A0 Valine--trn VARS
8,12107 0 P26641;P2 P26641;P2 Elongation EEF1G
2,64048 0,009186 P27105;F8 P27105 Erythrocyt STOM
10,5373 0 P27144;D3 P27144;D3 Adenylate AK4
7,64139 1,94E-05 P27338;P2 P27338;P2 Amine oxic MAOB
3,587 0,002792 P27348;E9 P27348;E9 14-3-3 pro YWHAQ
11,919 0 P27449;H3 P27449;H3 V-type pro ATP6V0C
10,1134 0 P27487;F8 P27487 Dipeptidyl DPP4
1,532 0,049407 X1WI28;P2 X1WI28;P2 60S riboso RPL10;RPL10L
9,66372 0 P27694;I3I P27694;I3I Replicator RPA1
13,2569 0 P27695;G3 P27695;G3 DNA-(apur APEX1
7,33845 3,6E-05 P27797;K7 P27797;K7 Calreticulir CALR
7,4527 2,76E-05 P27824;P2 P27824;P2 Calnexin CANX
4,14793 0,001393 X5D2R7;P2 X5D2R7;P2 Proteasom PSM8;PSMB8
2,84084 0,007165 P28066;P2 P28066;P2 Proteasom PSMA5
1,97687 0,024317 P28070 P28070 Proteasom PSMB4
0,950924 0,13202 P28072;A0 P28072;A0 Proteasom PSMB6
3,5381 0,002951 P28288;P2 P28288;P2 ATP-bindin ABCD3
8,82509 0 P28330 P28330 Long-chain ACADL
10,6101 0 P28331;P2 P28331;P2 NADH-ubic NDUFS1
3,61758 0,002692 P28482;P2 P28482;P2 Mitogen-a MAPK1
2,98563 0,005907 P28838-2;I P28838-2;I Cytosol am LAP3
6,90766 6,57E-05 P29401;P2 P29401;P2 Transketol TKT
3,59025 0,00278 P29590;P2 P29590;P2 Protein PV PML
1,4144 0,059794 P29992;K7 P29992;K7 Guanine ni GNA11
10,153 0 P30038;P3 P30038;P3 Delta-1-py ALDH4A1
18,3193 0 P30039;P3 P30039;P3 Phenazine PBLD
2,35596 0,013759 P30040;F8 P30040;F8 Endoplasm ERP29
5,07635 0,000479 P30041 P30041 Peroxiredc PRDX6
1,60202 0,044207 P30043;A0 P30043;A0 Flavin redu BLVRB
16,9365 0 P30044-2;I P30044-2;I Peroxiredc PRDX5
2,83666 0,007174 P30046;B5 P30046;B5 D-dopachr DDT;DDTL
10,6063 0 P30048;P3 P30048;P3 Thioredoxi PRDX3
8,13618 0 P30049 P30049 ATP synthz ATP5D
0,770709 0,183174 P30050;P3 P30050;P3 60S riboso RPL12
23,6358 0 P30084 P30084 Enoyl-CoA ECHS1
2,77264 0,007781 P30085;P3 P30085;P3 UMP-CMP CMPK1
8,65902 0 P30086 P30086 Phosphatic PEBP1
4,08221 0,001483 P30101 P30101 Protein dis PDIA3
5,6979 0,000291 P30153;B3 P30153;B3 Serine/thr PPP2R1A
14,2532 0 P30405;R4 P30405 Peptidyl-pr PPIF
3,18397 0,00455 P30480 P30480 HLA class I HLA-B
5,03796 0,000518 P30491 P30491 HLA class I HLA-B
1,86205 0,028848 P30536;B1 P30536;B1 Translocat TSPO
11,6274 0 P30613;P3 P30613;P3 Pyruvate k PKLR
2,66171 0,008988 P30711;A0 P30711;A0 Glutathion GSTT1
3,61369 0,002697 P30740;P3 P30740;P3 Leukocyte SERPINB1
8,05788 1,04E-05 P30837;A0 P30837 Aldehyde c ALDH1B1
21,532 0 P31040;P3 P31040;P3 Succinate c SDHA

8,8838 0 P31146;H3 P31146;H3 Coronin-1/ CORO1A
6,48035 0,000104 P31150;G5 P31150 Rab GDP d GDI1
4,86974 0,000592 P31153;P3 P31153;P3 S-adenosyl MAT2A
14,5567 0 P31639;P3 P31639;P3 Sodium/gli SLC5A2
1,26703 0,07659 P31689;P3 P31689;P3 DnaI homc DNAJA1
16,3627 0 P31930 P31930 Cytochrom UQCRC1
6,22607 0,000171 P31937;H7 P31937 3-hydroxyi HIBADH
4,19323 0,00128 P31939;P3 P31939;P3 Bifunctionα ATIC
4,60925 0,000794 P31942-3;I P31942-3;I Heterogen HNRNPH3
7,02331 5,88E-05 P31946-2;I P31946-2;I 14-3-3 pro YWHAB
2,04659 0,021869 P31948-3;I P31948-3;I Stress-indu STIP1
8,15526 0 P31949 P31949 Protein S1(S100A11
1,92757 0,026179 P32119;A6 P32119;A6 Peroxiredc PRDX2
3,87321 0,001959 P32189-1;I P32189-1;I Glycerol ki GK;GK3P
6,08383 0,000181 P32455;Q5 P32455 Interferon- GBP1
1,92421 0,026253 P32969;D6 P32969;D6 60S riboso RPL9
16,8268 0 P33121;E7 P33121;E7 Long-chain ACSL1
2,86538 0,006958 P33176 P33176 Kinesin-1 h KIF5B
6,32612 0,000152 P33316;HC P33316;HC Deoxyuridi DUT
25,8963 0 P34896-2;I P34896-2;I Serine hyd SHMT1
5,43459 0,00041 P34897-3;I P34897-3;I Serine hyd SHMT2
6,99808 5,85E-05 P34913;E5 P34913;E5 Bifunctionα EPHX2
1,28481 0,074239 P34932;A0 P34932;A0 Heat shock HSPA4
7,69897 1,96E-05 P34949;P3 P34949;P3 Mannose-ε MPI
8,84974 0 P35221;P3 P35221;P3 Catenin αI CTNNA1
11,3728 0 P35232;C9 P35232;C9 Prohibitin PHB
1,14849 0,093146 P35270;DI P35270 Sepiapterir SPR
6,86315 6,52E-05 P35555;HC P35555 Fibrillin-1 FBN1
7,9041 2,05E-05 P35558;P3 P35558 Phosphoer PCK1
1,48606 0,053157 P35573-2;I P35573-2;I Glycogen c AGL
10,911 0 P35579;Q5 P35579 Myosin-9 MYH9
4,79984 0,000624 P35580-2;I P35580-2;I Myosin-10 MYH10
5,81055 0,000261 P35606;P3 P35606;P3 Coatomer COPB2
8,17314 0 P35749-4;I P35749-4;I Myosin-11 MYH11
5,20237 0,000452 P35754 P35754 Glutaredo GLRX
1,32099 0,070088 P35914;P3 P35914;P3 Hydroxymε HMGCL
0,132266 0,683456 P35998;P3 P35998;P3 26S protea PSMC2
6,96564 6,64E-05 P36269;P3 P36269;P3 Gamma-gli GGT5
14,7278 0 P36542;P3 P36542;P3 ATP synthε ATP5C1
18,5336 0 P36543;P3 P36543;P3 V-type pro ATP6V1E1
0,888023 0,147504 P36578;H3 P36578;H3 60S riboso RPL4
13,7721 0 P36776;K7 P36776;K7 Lon protea LONP1
2,68603 0,008746 P36871;A0 P36871;A0 Phosphogli PGM1
4,17659 0,001316 P36955;I3I P36955 Pigment εI SERPINF1
11,0936 0 P36957;P3 P36957;P3 Dihydrolipi DLST
5,62832 0,000316 P37235;P8 P37235;P8 Hippocalcii HPCAL1;HPCA;NCALD
12,6744 0 P37802;X6 P37802;X6 Transgelin- TAGLN2
1,10445 0,100835 P37837;F2 P37837;F2 Transaldoli TALDO1
6,97169 6,65E-05 P38117;P3 P38117;P3 Electron tr ETFB
24,0883 0 P38606;P3 P38606;P3 V-type pro ATP6V1A

17,3742 0 P38646;A0 P38646 Stress-70 ç HSPA9
0,53267 0,288518 P39023;G5 P39023;G5 60S riboso RPL3
6,54684 9,21E-05 P39060-2;I P39060-2;I Collagen al COL18A1
7,73188 1,97E-05 P40121;P4 P40121;P4 Macrophag CAPG
1,86366 0,028778 P40227;P4 P40227;P4 T-complex CCT6A
9,13866 0 P40261 P40261 Nicotinami NNMT
11,3021 0 P40306;J3I P40306;J3I Proteasom PSMB10
0,342945 0,423792 P40429;Ml P40429;Ml 60S riboso RPL13A;RPL13a
6,94744 6,61E-05 P40763-3;I P40763-3;I Signal tran STAT3
15,2377 0 P40925;P4 P40925;P4 Malate del MDH1
13,796 0 P40926;P4 P40926;P4 Malate del MDH2
11,7643 0 P40939;HC P40939;HC Trifunction HADHA
1,15018 0,092976 P41091;Q2 P41091;Q2 Eukaryotic EIF2S3;EIF2S3L
2,17372 0,017734 P41250;DI P41250 Glycine--tR GARS
6,78545 7,26E-05 P42126;Q5 P42126;Q5 Enoyl-CoA EC11;DCI
1,85123 0,029378 P42167-2;I P42167-2;I Lamina-ass TMPO
7,85061 2,02E-05 P42224;P4 P42224;P4 Signal tran STAT1
10,28 0 P42704;A0 P42704 Leucine-ric LRPPRC
9,84329 0 P42765;A0 P42765;A0 3-ketoacyl- ACAA2
3,8414 0,002019 P42785;P4 P42785;P4 Lysosomal PRCP
3,20186 0,004473 P43034;I3I P43034 Platelet-ac PAFAH1B1
5,56207 0,000351 P43121;P4 P43121;P4 Cell surfac MCAM
10,3037 0 P43155-2;I P43155-2;I Carnitine C CRAT
5,57534 0,000332 P43304;P4 P43304;P4 Glycerol-3- GPD2
7,18774 4,34E-05 P43490;A0 P43490;A0 Nicotinami NAMPT
1,58879 0,045119 P43652 P43652 Afamin AFM
1,01166 0,119189 P43686-2;I P43686-2;I 26S protea PSMC4
11,2438 0 P43897;P4 P43897;P4 Elongation TSFM
17,4908 0 P45880;A0 P45880;A0 Voltage-de VDAC2
12,9357 0 P45954;P4 P45954;P4 Short/bran ACADSB
4,19073 0,001279 P45974-2;I P45974-2;I Ubiquitin c USP5
11,8584 0 P46063;F8 P46063 ATP-depen RECQL
4,13599 0,001401 P46939;P4 P46939;P4 Utrophin UTRN
5,69786 0,000291 P46940;A0 P46940;A0 Ras GTPasç IQGAP1
2,17576 0,017697 P46952;C9 P46952;C9 3-hydroxyç HAAO
6,68711 7,07E-05 P47755;F8 P47755;F8 F-actin-cap CAPZA2
4,05367 0,001544 P47756-2;I P47756-2;I F-actin-cap CAPZB
0,647117 0,230309 P47897;P4 P47897;P4 Glutamine- QARS
4,88414 0,00059 P47985;P0 P47985;P0 Cytochrom UQCRFS1;UQCRFS1P1
11,1765 0 P48047;A0 P48047;A0 ATP synthç ATP5O
2,64421 0,009154 P48059-2;I P48059-2;I LIM and se LIMS1
3,4208 0,00347 P48637;A0 P48637;A0 Glutathion GSS
2,74671 0,008058 P48643;B7 P48643;B7 T-complex CCT5
16,9224 0 P48735;P4 P48735;P4 Isocitrate c IDH2
5,52581 0,000374 P49189;P4 P49189;P4 4-trimethy ALDH9A1
2,22007 0,016643 P49257 P49257 Protein ER LMAN1
0,774949 0,181918 P49406;A0 P49406;A0 39S riboso MRPL19
19,5552 0 P49411;H3 P49411 Elongation TUFM
6,10372 0,000182 P49419-2;I P49419-2;I Alpha-amir ALDH7A1
0,295441 0,469291 P49588;P4 P49588;P4 Alanine--tF AARS

3,40424 0,003511 Q5T5C7;P4 Q5T5C7;P4 Serine--trf SARS
2,25242 0,015883 P49720;A0 P49720;A0 Proteasom PSMB3
6,47545 0,000104 P49721;A0 P49721;A0 Proteasom PSMB2
7,1104 4,27E-05 P49748-3;I P49748-3;I Very long-c ACADVL
8,16731 0 P49755;G3 P49755 Transmem TMED10
2,82105 0,007308 P49773;D6 P49773;D6 Histidine tr HINT1
1,45148 0,056352 P49915;P4 P49915;P4 GMP synth GMPS
12,4435 0 P50053 P50053 Ketohexok KHK
6,688 7,09E-05 P50135;P5 P50135;P5 Histamine HNMT
12,3357 0 P50213;HC P50213;HC Isocitrate c IDH3A
6,28664 0,000158 P50395;P5 P50395;P5 Rab GDP d GDI2
2,56711 0,010182 Q5HY57;P5 Q5HY57;P5 Emerin EMD
11,8248 0 P50416;P5 P50416;P5 Carnitine C CPT1A
16,7395 0 P50440-2;I P50440-2;I Glycine am GATM
2,52946 0,010818 P50453;P5 P50453 Serpin B9 SERPINB9
7,49994 2,78E-05 P50454;E9 P50454;E9 Serpin H1 SERPINH1
1,02389 0,116774 Q3KNR6;P1 Q3KNR6;P1 Hsc70-inte ST13;ST13P5;ST13P4
2,3995 0,012941 P50570-2;I P50570-2;I Dynamin-2 DNM2
6,37861 0,000139 P50990;P5 P50990;P5 T-complex CCT8
6,51313 0,000106 P50991;P5 P50991;P5 T-complex CCT4
9,36296 0 P50995-2;I P50995-2;I Annexin A1 ANXA11
0,723386 0,200014 P51148;P5 P51148;P5 Ras-relatec RAB5C
15,4987 0 P51149;C9 P51149;C9 Ras-relatec RAB7A
14,9181 0 P51553;P5 P51553;P5 Isocitrate c IDH3G
8,18557 0 P51570;P5 P51570;P5 Galactokin GALK1
1,21318 0,083721 P51571;A6 P51571;A6 Translocon SSR4
1,50772 0,051326 P51572;P5 P51572;P5 B-cell rece| BCAP31
8,20135 0 P51648-2;I P51648-2;I Fatty alde| ALDH3A2
8,90092 0 P51649-2;I P51649-2;I Succinate-c| ALDH5A1
2,65018 0,009096 P51665;H3 P51665;H3 26S protea PSMD7
8,00639 1,03E-05 P51687;DI P51687 Sulfite ox|c SUOX
7,58573 1,89E-05 P51884 P51884 Lumican LUM
3,75429 0,002296 P51888 P51888 Prolargin PRELP
3,63105 0,002652 P51911;P5 P51911;P5 Calponin-1 CNN1
6,57327 9,27E-05 P51991;P5 P51991;P5 Heterogen HNRNPA3
4,55993 0,000839 P52209;P5 P52209;P5 6-phospho PGD
5,08653 0,000475 P52272;A0 P52272;A0 Heterogen HNRNPM
9,93874 0 P52566;HC P52566;HC Rho GDP-d ARHGDIB
1,50149 0,05181 P52597;A0 P52597 Heterogen HNRNPF
11,9159 0 P52758;HC P52758;HC Ribonuclea HRSP12
1,18581 0,087598 P52788;P5 P52788;P5 Spermine s SMS
8,31023 0 P52907 P52907 F-actin-cap CAPZA1
3,47697 0,003222 P53004;C9 P53004 Biliverdin r BLVRA
0,813587 0,169682 P53007;B4 P53007;B4 Tricarboxyl SLC25A1
14,7686 0 P53396-2;I P53396-2;I ATP-citrate ACLY
19,333 0 P53597;A0 P53597;A0 Succinyl-Cc SUCLG1
0,975703 0,12636 P53618;E9 P53618;E9 Coatomer : COPB1
4,14736 0,001391 P53621;P5 P53621;P5 Coatomer : COPA
15,0729 0 P53634;HC P53634 Dipeptidyl CTSC
2,21628 0,016709 P53992;G5 P53992;G5 Protein tra SEC24C

0,825038 0,166012 P54136-2;I P54136-2;I Arginine--t RARS
1,64058 0,041336 P54802 P54802 Alpha-N-ac NAGLU
1,00792 0,11984 P54920;M P54920;M Alpha-solu NAPA
0,905115 0,142918 P55060-3;I P55060-3;I Exportin-2 CSE1L
2,62533 0,0094 P55072;C9 P55072 Transitionε VCP
10,7522 0 P55084-2;I P55084-2;I Trifunction HADHB
16,8849 0 P55268;F5 P55268 Laminin su LAMB2
6,14527 0,000177 P55290;P5 P55290;P5 Cadherin-1 CDH13
0,607102 0,248986 P55735;A8 P55735;A8 Protein SE(SEC13
2,04069 0,022074 P55795 P55795 Heterogen HNRNPH2
9,69683 0 P55809;E9 P55809;E9 Succinyl-Cc OXCT1
1,94 0,025672 P55884;P5 P55884;P5 Eukaryotic EIF3B
6,00999 0,000213 P56134;P5 P56134;P5 ATP synthε ATP5J2
7,51715 2,8E-05 P56134-2;I P56134-2;I ATP synthε ATP5J2;ATP5J2-PTCD1
5,0217 0,000514 P56192;P5 P56192;P5 Methionin MARS
9,81942 0 P56385 P56385 ATP synthε ATP5I
0,283507 0,482195 P56470;M P56470;M Galectin-4; LGALS4
3,83533 0,002031 P56537;P5 P56537;P5 Eukaryotic EIF6
0,984875 0,124511 P57105;A0 P57105;A0 Synaptojar SYNJ2BP;SYNJ2BP-COX16
3,61612 0,002691 P58546;C9 P58546;C9 Myotrophi MTPN
2,68226 0,008779 P60174-1;I P60174-1;I Triosephos TPI1
2,53889 0,010692 P60228;E5 P60228;E5 Eukaryotic EIF3E
7,22191 4,37E-05 P60660;F8 P60660;F8 Myosin ligl MYL6
1,75905 0,034025 P60709;A0 P60709;A0 Actin, cyto ACTB
8,0817 1,04E-05 P60842;P6 P60842;P6 Eukaryotic EIF4A1
1,51121 0,051091 P60900;G3 P60900;G3 Proteasom PSMA6
7,62668 1,93E-05 P60903 P60903 Protein S1(S100A10
0,846968 0,159653 P60953;P6 P60953;P6 Cell divisio CDC42
3,86979 0,001962 P60981;P6 P60981;P6 Destrin DSTN
0,78057 0,180132 P61006;P6 P61006;P6 Ras-relateε RAB8A
9,70394 0 P61019;P6 P61019;P6 Ras-relateε RAB2A
4,32413 0,001121 P61026 P61026 Ras-relateε RAB10
4,38351 0,00107 P61077;P6 P61077;P6 Ubiquitin-c UBE2D3
4,06085 0,001535 P61088;F8 P61088;F8 Ubiquitin-c UBE2N
7,07106 5,08E-05 P61106;X6 P61106;X6 Ras-relateε RAB14
2,81144 0,007427 P61158;B4 P61158;B4 Actin-relat ACTR3
2,29795 0,014837 P61160;P6 P61160;P6 Actin-relat ACTR2
0,016743 0,943826 P61163;R4 P61163;R4 Alpha-cent ACTR1A
1,45726 0,055796 P61224;P6 P61224;P6 Ras-relateε RAP1B
6,20324 0,000178 P61313;E7 P61313;E7 60S riboso RPL15
0,945717 0,133073 P61457;DI P61457 Pterin-4-al PCBD1
0,249216 0,520204 P61586;C9 P61586;C9 Transformi RHOA
4,82017 0,000604 P61604;B8 P61604;B8 10 kDa heε HSPE1
11,7441 0 P61626;F8 P61626;F8 Lysozyme (LYZ
1,39598 0,06179 P61769;HC P61769;HC Beta-2-mic B2M
0,173561 0,61858 P61970;H3 P61970;H3 Nuclear trε NUTF2
6,48926 0,000105 P61978-2;I P61978-2;I Heterogen HNRNPK
6,0237 0,0002 P61981 P61981 14-3-3 pro YWHAG
0,125164 0,695474 P62081;B5 P62081;B5 40S riboso RPS7
3,63825 0,00263 P62140;E7 P62140;E7 Serine/thrε PPP1CB

1,58109 0,045548 P62195;P6 P62195;P6 26S protea PSMC5
1,56239 0,047168 P62241;Q5 P62241;Q5 40S riboso RPS8
0,173567 0,618984 P62244;I3I P62244;I3I 40S riboso RPS15A
0,474371 0,32238 P62249;MI P62249;MI 40S riboso RPS16
3,11625 0,004996 P62258;P6 P62258;P6 14-3-3 pro YWHAE
0,698662 0,208815 P62269;A0 P62269 40S riboso RPS18
6,58299 8,51E-05 P62306;A0 P62306 Small nuclε SNRPF
2,49435 0,01132 P62314;J3I P62314;J3I Small nuclε SNRPD1
2,72656 0,008298 P62318;P6 P62318;P6 Small nuclε SNRPD3
4,04138 0,001568 P62491;H3 P62491;H3 Ras-relatec RAB11A
1,714 0,036641 P62701;Q8 P62701 40S riboso RPS4X
4,71112 0,000691 P62736;P6 P62736;P6 Actin, aorti ACTA2;ACTG2
0,484891 0,316433 P62753;A2 P62753;A2 40S riboso RPS6
0,515587 0,29808 P62805 P62805 Histone H4 HIST1H4A
5,95379 0,000231 P62820;E7 P62820;E7 Ras-relatec RAB1A
1,65662 0,040218 P62834;A0 P62834;A0 Ras-relatec RAP1A
7,19211 4,35E-05 P62873;P6 P62873;P6 Guanine nι GNB1
5,43307 0,000409 P62879;E7 P62879;E7 Guanine nι GNB2
0,55585 0,275619 P62906 P62906 60S riboso RPL10A
0,803757 0,172566 P62913;Q5 P62913;Q5 60S riboso RPL11
3,68207 0,002489 P62937;P6 P62937;P6 Peptidyl-pr PPIA
5,12349 0,00047 P62942;Q1 P62942;Q1 Peptidyl-pr FKBP1A;FKBP12-Exin;FKBP12-Exip2
2,91192 0,006543 P62979;P0 P62979;P0 Ubiquitin-4 RPS27A;UBC;UBB;UBA52
14,3648 0 P62993;P6 P62993;P6 Growth fac GRB2
4,52385 0,000877 P62995;P6 P62995;P6 Transformι TRA2B
0,575805 0,264765 P63000;P6 P63000;P6 Ras-relatec RAC1;RAC3
2,13437 0,018927 P63010-3;I P63010-3;I AP-2 comp AP2B1
0,304323 0,460962 P63092-3;I P63092-3;I Guanine nι GNAS
7,11526 4,28E-05 P63096;P6 P63096;P6 Guanine nι GNAI1
8,96815 0 P63104;E7 P63104;E7 14-3-3 pro YWHAZ
4,23709 0,001232 P63151;P6 P63151;P6 Serine/thrε PPP2R2A
4,38117 0,001068 P63172 P63172 Dynein ligι DYNLT1
0,486814 0,315344 Q8WVC2;F Q8WVC2;F 40S riboso RPS21
5,85853 0,000249 P63241;I3I P63241;I3I Eukaryotic EIF5A;EIF5AL1;EIF5A2
4,36734 0,001072 P63244;J3I P63244;J3I Guanine nι GNB2L1
7,36139 2,73E-05 P63261;I3I P63261 Actin, cyto ACTG1
1,17952 0,088504 P67775;P6 P67775;P6 Serine/thrε PPP2CA
4,84567 0,000595 P67936;A0 P67936;A0 Tropomyo: TPM4
6,71575 7,11E-05 P68032 P68032 Actin, alph ACTC1
2,24706 0,016 Q5VTE0;P6 Q5VTE0;P6 Putative el EEF1A1P5;EEF1A1
13,7579 0 P68363;P6 P68363;P6 Tubulin alι TUBA1B
4,6045 0,000798 P68366-2;I P68366-2;I Tubulin alι TUBA4A
7,35824 2,72E-05 P68371;MI P68371 Tubulin be TUBB4B
0,388266 0,385098 P68402;P6 P68402;P6 Platelet-ac PAFAH1B2
8,75432 0 P68431 P68431 Histone H3 HIST1H3A
2,13329 0,018935 P68871;A0 P68871;A0 Hemoglobi HBB
2,3823 0,013259 P69905;A0 P69905;A0 Hemoglobi HBA1
7,61759 1,92E-05 P78371;P7 P78371;P7 T-complex CCT2
3,70315 0,002413 P78417;Q5 P78417;Q5 Glutathion GSTO1
11,3165 0 P78527;P7 P78527;P7 DNA-depei PRKDC

15,6869	0	P80365;DI	P80365	Corticoster	HSD11B2
5,15135	0,000474	P82980;A0	P82980;A0	Retinol-bin	RBP5
5,19809	0,000451	P84077;F5	P84077;F5	ADP-ribosy	ARF1
3,04393	0,005484	P84085;C9	P84085;C9	ADP-ribosy	ARF5
12,2865	0	P84090;G3	P84090;G3	Enhancer c	ERH
10,9406	0	P98082-2;I	P98082-2;I	Disabled h	DAB2
1,54283	0,048662	P98095;P9	P98095;P9	Fibulin-2	FBLN2
6,44808	0,000111	P98160;A0	P98160	Basement	HSPG2
20,0075	0	P98164;A0	P98164	Low-densit	LRP2
21,243	0	Q00325-2;	Q00325-2;	Phosphate	SLC25A3
10,8864	0	Q00796;H	Q00796;H	Sorbitol de	SORD
3,61194	0,002698	Q00839;Q	Q00839;Q	Heterogen	HNRNPU
0,996062	0,122094	Q01082;A	Q01082;A	Spectrin b	SPTBN1
6,59222	8,58E-05	Q01432-6;	Q01432-6;	AMP deam	AMPD3
6,14305	0,000177	Q01518;Q	Q01518	Adenylyl c	CAP1
9,19492	0	Q01740;Q	Q01740;Q	Dimethyla	FMO1
14,4222	0	Q01813;Q	Q01813;Q	ATP-depen	PFKP
5,58101	0,000333	Q01995;H	Q01995;H	Transgelin	TAGLN
18,3194	0	Q02218;E	Q02218;E	2-oxogluta	OGDH
7,61179	1,91E-05	Q02252;Q	Q02252;Q	Methylmal	ALDH6A1
9,9472	0	Q02338;E	Q02338;E	D-beta-hyc	BDH1
2,51185	0,01104	Q02543;M	Q02543;M	60S riboso	RPL18A
3,38808	0,003556	Q02878;F	Q02878	60S riboso	RPL6
14,0277	0	Q02928;A	Q02928;A	Cytochrom	CYP4A11
6,09383	0,000182	Q02952-3;	Q02952-3;	A-kinase a	AKAP12
12,6629	0	Q02978;I	Q02978;I	Mitochond	SLC25A11
11,7914	0	Q03154;Q	Q03154;Q	Aminoacyl	ACY1;ABHD14A-ACY1
4,82281	0,000605	Q03252;D	Q03252	Lamin-B2	LMNB2
3,36609	0,003622	Q03591;B	Q03591;B	Compleme	CFHR1
7,89476	2,04E-05	Q04446;E	Q04446;E	1,4-alpha-	GBE1
0,566591	0,269526	Q04760;Q	Q04760;Q	Lactoylglut	GLO1
10,6126	0	Q04828;H	Q04828;H	Aldo-keto	AKR1C1
15,2122	0	Q04837;A	Q04837;A	Single-stra	SSBP1
9,60966	0	Q04917;A	Q04917;A	14-3-3 pro	YWHAH
4,88212	0,000589	Q05655;Q	Q05655;Q	Protein kin	PRKCD
8,69382	0	Q05682-4;	Q05682-4;	Caldesmor	CALD1
7,87175	2,03E-05	Q05707;Q	Q05707;Q	Collagen al	COL14A1
2,09241	0,020333	Q06033-2;	Q06033-2;	Inter-alpha	ITIH3
2,33056	0,014156	Q06210-2;	Q06210-2;	Glutamine	GFPT1
3,43746	0,003402	Q06323;Q	Q06323;Q	Proteasom	PSME1
14,7804	0	Q06830;A	Q06830;A	Peroxiredc	PRDX1
3,75337	0,002298	Q07021;I	Q07021;I	Compleme	C1QBP
11,3396	0	Q07065	Q07065	Cytoskelet	CKAP4
3,82376	0,002068	Q07075	Q07075	Glutamyl a	ENPEP
4,95653	0,000548	Q07507	Q07507	Dermatop	DPT
1,22043	0,082772	Q07954;H	Q07954	Prolow-dei	LRP1
9,79423	0	Q07960;H	Q07960;H	Rho GTPas	ARHGAP1
4,50683	0,00088	Q08211;Q	Q08211	ATP-depen	DHX9
5,68084	0,00029	Q08257;A	Q08257;A	Quinone o	CRYZ
4,50446	0,000879	Q08380;K	Q08380	Galectin-3-	LGALS3BP

8,36956 0 Q08426;Q(Q08426;Q(Peroxisom EHHADH
10,6864 0 Q08AH3;F(Q08AH3;F(Acyl-coenz ACSM2A
0,425073 0,356329 Q09028-3; Q09028-3; Histone-bii RBBP4;RBBP7
15,687 0 Q09666;E(Q09666 Neuroblast AHNAK
2,15149 0,018431 Q10567-3; Q10567-3; AP-1 comp AP1B1
4,86559 0,000591 Q10713;Q(Q10713;Q(Mitochond PMPCA
14,2287 0 Q12765;Q(Q12765;Q(Secernin-1 SCR1
5,98378 0,000218 Q12792;Q(Q12792;Q(Twinfilin-1 TWF1
3,69803 0,002421 Q12805-2; Q12805-2; EGF-contai EFEMP1
1,70067 0,037422 Q12874 Q12874 Splicing fac SF3A3
7,20379 4,36E-05 Q12905;B(Q12905;B(Interleukin ILF2
1,88968 0,027624 Q12906-5; Q12906-5; Interleukin ILF3
6,53181 0,000107 Q12907;D(Q12907;D(Vesicular ir LMAN2
7,36895 2,74E-05 Q12931-2; Q12931-2; Heat shock TRAP1
7,5038 2,79E-05 Q13011;M Q13011;M Delta(3,5)- ECH1
2,33654 0,014079 Q13045-2; Q13045-2; Protein flig FLII
1,39014 0,062351 Q13057;Q(Q13057;Q(Bifunction(COASY
1,84305 0,029845 Q13131-2; Q13131-2; 5-AMP-act PRKAA1
0,812335 0,170009 Q13148;A(Q13148;A(TAR DNA-t TARDBP;TDP43
0,480796 0,31901 Q13155;A(Q13155;A(Aminoacyl AIMP2
3,02534 0,005596 Q13177;H(Q13177 Serine/thr(PAK2
1,6114 0,043462 Q13185;S4 Q13185;S4 Chromobo CBX3
0,960704 0,129862 Q13200;Q(Q13200;Q(26S protea PSMD2
4,41376 0,001029 Q13228-4; Q13228-4; Selenium-k SELENBP1
2,34748 0,013885 Q13232;H(Q13232;H(Nucleoside NME3
0,090878 0,761474 Q13247-3; Q13247-3; Serine/argi SRSF6
2,21555 0,016715 Q13263-2; Q13263-2; Transcripti TRIM28
2,641 0,009187 Q13347;Q(Q13347 Eukaryotic EIF3I
0,253058 0,515879 Q13404;G(Q13404;G(Ubiquitin-c UBE2V1;TMEM189-UBE2V1
3,76592 0,002256 Q13409-2; Q13409-2; Cytoplasm DYNC1I2
24,5808 0 Q13423;E(Q13423;E(NAD(P) tra NNT
1,50688 0,051346 Q13425;A(Q13425 Beta-2-syn SNTB2
2,49439 0,011331 Q13492;Q(Q13492;Q(Phosphatic PICALM
2,28744 0,01508 Q13505-3; Q13505-3; Metaxin-1 MTX1
6,88988 6,56E-05 Q13509;A(Q13509;A(Tubulin be TUBB3
1,69917 0,037442 Q13561;Q(Q13561;Q(Dynactin si DCTN2
9,1094 0 Q13576;F5 Q13576;F5 Ras GTPas(IQGAP2
18,5023 0 Q13642-1; Q13642-1; Four and a FHL1
0,961765 0,129682 Q13724-2; Q13724-2; Mannosyl- MOGS
3,44009 0,003406 Q13813;Q(Q13813;Q(Spectrin al SPTAN1
21,1404 0 Q13825;Q(Q13825;Q(Methylglut AUH
10,7069 0 Q13838-2; Q13838-2; Spliceosonr DDX39B
6,51268 0,000106 Q13884;A(Q13884;A(Beta-1-syn SNTB1
9,58397 0 Q14019;H(Q14019;H(Coactosin- COTL1
3,34668 0,003679 Q14103-3; Q14103-3; Heterogen HNRNPD
5,93157 0,000237 Q14112;A(Q14112;A(Nidogen-2 NID2
8,55065 0 Q14117;E5 Q14117 Dihydropy(DPYS
1,36112 0,065491 Q14152-2; Q14152-2; Eukaryotic EIF3A
5,49558 0,000382 Q14165;H(Q14165 Malectin MLEC
3,69905 0,002424 Q14166 Q14166 Tubulin--ty TLL12

3,76781 0,002259 Q14195;Q: Q14195;Q: Dihydropyridyl DPYSL3
3,11549 0,004991 Q14204;A: Q14204;A: Cytoplasm DYNC1H1
0,741069 0,193303 Q14247-3; Q14247-3; Src substratum CTTN
4,04729 0,001555 Q14353;A: Q14353;A: Guanidino: GAMT
1,08323 0,104734 Q14376;Q: Q14376;Q: UDP-glucose GALE
1,99896 0,023497 Q14444-2; Q14444-2; Caprin-1 CAPRIN1
5,99752 0,000212 Q14651;C: Q14651 Plastin-1 PLS1
15,4345 0 Q14697;F5 Q14697;F5 Neutral alpha GANAB
11,7791 0 Q14764;H: Q14764 Major vault MVP
8,36156 0 Q14847;Q: Q14847;Q: LIM and SH LASP1
8,17106 0 Q14894;I3 Q14894;I3 Ketimine rich CRYM
7,45215 2,75E-05 Q14974;Q: Q14974;Q: Importin subunit KPNB1
5,18738 0,000449 Q14980-2; Q14980-2; Nuclear matrix NUMA1
7,59919 1,9E-05 Q15019-2; Q15019-2; Septin-2 SEPT2
3,24432 0,004236 Q15029-2; Q15029-2; 116 kDa UBEFTUD2
9,92223 0 Q15031;A: Q15031;A: Probable kinase LARS2
0,630697 0,238065 Q15046;Q: Q15046;Q: Lysine--transferase KARS
2,06801 0,021093 Q15063-3; Q15063-3; Periostin POSTN
3,24874 0,004215 Q15067-2; Q15067-2; Peroxisomal ACOX1
9,88228 0 Q15084-3; Q15084-3; Protein disulfide PDIA6
0,991248 0,123088 Q15124;Q: Q15124;Q: Phosphoglucose PGM5
12,6444 0 Q15149-7; Q15149-7; Plectin PLEC
0,301565 0,463582 Q15181;Q: Q15181;Q: Inorganic pyrophosphate PPA1
0,160116 0,638751 Q15233-2; Q15233-2; Non-POU class NONO
9,27777 0 Q15274;C: Q15274;C: Nicotinate: quinone QPRT
5,3221 0,00043 Q15365 Q15365 Poly(rC)-binding PCBP1
0,959787 0,129924 Q15366-4; Q15366-4; Poly(rC)-binding PCBP2
2,74845 0,008039 Q15393;Q: Q15393 Splicing factor SF3B3
7,59509 1,9E-05 Q15404;Q: Q15404;Q: Ras suppressor RSU1
3,80422 0,002141 Q15417;Q: Q15417;Q: Calponin-3 CNN3
4,13066 0,001402 Q15582;H: Q15582;H: Transforming TGFBI
3,78006 0,002197 Q15654;H: Q15654;H: Thyroid receptor TRIP6
1,2362 0,080729 Q15717;Q: Q15717;Q: ELAV-like factor ELAVL1
3,9723 0,001668 Q15746-4; Q15746-4; Myosin light MYLK
13,7883 0 Q15907;Q: Q15907;Q: Ras-related RAB11B
14,6855 0 Q16134;Q: Q16134;Q: Electron transfer ETFDH
1,31573 0,070596 Q16401-2; Q16401-2; 26S proteasome PSMD5
5,5019 0,000383 Q16531;F5 Q16531;F5 DNA damage DDB1
1,93527 0,025826 Q16543;K7 Q16543;K7 Hsp90 co-chaperone CDC37
5,03105 0,000515 Q16658;C: Q16658 Fascin FSCN1
14,6137 0 Q16698-2; Q16698-2; 2,4-dienoyl DECR1
1,71245 0,036779 Q16718;A: Q16718;A: NADH dehydrogenase NDUFA5
7,23136 4,39E-05 Q16762;B1 Q16762 Thiosulfate TST
0,527555 0,291419 Q16777;Q: Q16777;Q: Histone H2 HIST2H2AC;HIST2H2AA3
11,7573 0 Q16790;A: Q16790;A: Carbonic anhydrase CA9
6,14519 0,000177 Q16795;A: Q16795;A: NADH dehydrogenase NDUFA9
14,8243 0 Q16798;E: Q16798;E: NADP-dependent ME3
12,571 0 Q16822;Q: Q16822;Q: Phosphoenolpyruvate PCK2
17,047 0 Q16836;A: Q16836;A: Hydroxyacyl HADH
6,73218 7,13E-05 Q16853;Q: Q16853;Q: Membrane AOC3

5,74148 0,000286 Q16864;Q: Q16864;Q: V-type pro ATP6V1F
16,1234 0 Q16891-2; Q16891-2; MICOS conIMMT
4,05668 0,001546 Q27J81-2;Q: Q27J81-2;Q: Inverted fc INF2
13,067 0 Q3LXA3;H: Q3LXA3;H: Bifunctional DAK;TKFC
6,87365 6,53E-05 Q3ZCM7;A: Q3ZCM7;A: Tubulin beta TUBB8
3,65221 0,002606 Q3ZCQ8;Q: Q3ZCQ8;Q: Mitochondrial TIMM50
2,03212 0,022375 Q53GQ0;A: Q53GQ0;A: Very-long-chain HSD17B12
2,6467 0,009136 Q53H82 Q53H82 Beta-lactar LACTB2
9,08048 0 Q56VL3;J3: Q56VL3 OCIA domain OCIAD2
2,87094 0,006925 Q5JRX3;Q5: Q5JRX3;Q5: Presequence PITRM1
0,680509 0,215828 X6R3N0;Q: X6R3N0;Q: Long-chain SLC27A3
5,33254 0,000431 Q5R3I4;A9: Q5R3I4;A9: Tetratricolporin TTC38
2,30975 0,014549 Q5S007;E9: Q5S007 Leucine-rich LRRK2
0,336365 0,429926 Q5SSJ5-2;C: Q5SSJ5-2;C: Heterochromatin HP1BP3
18,0149 0 Q5T2W1;C: Q5T2W1;C: Na(+)/H(+) PDZK1;PDZK1P1
8,83114 0 Q5T440;DI: Q5T440 Putative transmembrane IBA57
0,947806 0,132741 Q5T6H7;Q: Q5T6H7;Q: Xaa-Pro an XPNPEP1
2,53961 0,010679 Q5T8A0;Q: Q5T8A0;Q: 28S ribosomal MRPS2
0,113819 0,716329 Q5TEC6 Q5TEC6 Histone H3 HIST2H3PS2
11,7336 0 Q68CK6;H: Q68CK6;H: Acyl-coenzyme ACSM2B
1,46744 0,054796 Q6DD88;F: Q6DD88;F: Atlantin-3 ATL3
14,9521 0 Q6IB77;Q6: Q6IB77 Glycine N-acyl GLYAT
2,58711 0,009938 Q6ICL3-3;C: Q6ICL3-3;C: Transporter TANGO2
1,26568 0,076746 Q6NUK1;Q: Q6NUK1;Q: Calcium-binding SLC25A24
6,49422 0,000105 Q6NVY1;Q: Q6NVY1;Q: 3-hydroxy HIBCH
7,91882 2,05E-05 Q6NZI2;Q6: Q6NZI2;Q6: Polymerase PTRF
10,4785 0 Q6P2Q9;I3: Q6P2Q9 Pre-mRNA PRPF8
6,43514 0,000118 Q6P4A8 Q6P4A8 Phospholipase PLBD1
2,08895 0,020447 Q6P587;Q: Q6P587;Q: Acylpyruvate FAHD1
14,3866 0 Q6PCB0;J3: Q6PCB0 von Willebrand VWA1
7,98298 1,03E-05 Q6XQN6;C: Q6XQN6;C: Nicotinate NAPRT
10,8248 0 Q6YN16;Q: Q6YN16;Q: Hydroxysteroid HSDL2
1,40598 0,060745 Q6ZUJ8;Q6: Q6ZUJ8;Q6: Phosphoinositide PIK3AP1
4,05836 0,001548 Q709F0;Q: Q709F0;Q: Acyl-CoA dehydrogenase ACAD11
7,29331 3,56E-05 Q71DI3 Q71DI3 Histone H3 HIST2H3A
12,1372 0 Q71U36;Q: Q71U36;Q: Tubulin alpha TUBA1A
7,17699 4,32E-05 Q7KZF4;H7: Q7KZF4 Staphylococcal SND1
1,52534 0,049908 Q7L266;Q: Q7L266;Q: Isoaspartyl transferase ASRGL1
5,04015 0,000513 Q7L576;Q: Q7L576;Q: Cytoplasmic CYFIP1
2,52311 0,010895 Q7Z3D6-2; Q7Z3D6-2; UPF0317 protein C14orf159
3,53928 0,00295 Q7Z406;Q: Q7Z406;Q: Myosin-14 MYH14
9,13289 0 Q7Z4W1;J: Q7Z4W1;J: L-xylulose 5-epimerase DCXR
0,623649 0,241151 Q7Z6Z7-2;Q: Q7Z6Z7-2;Q: E3 ubiquitin HUWE1
2,52051 0,010926 Q7Z7H5-3; Q7Z7H5-3; Transmembrane TMED4
5,44946 0,000405 Q86UP2-2; Q86UP2-2; Kinectin KTN1
1,55523 0,047721 Q86UX7-2; Q86UX7-2; Fermitin family FERMT3
6,28042 0,000158 Q86VP6;Q: Q86VP6;Q: Cullin-associated CAND1
12,296 0 Q86VU5;R: Q86VU5;R: Catechol O-methyltransferase COMTD1
8,88313 0 Q86WA6-2 Q86WA6-2 Valacyclovir BPHL
5,93489 0,000231 Q86WU2-2 Q86WU2-2 Probable DNA LDHD

8,41609 0 Q86XE5;Q Q86XE5;Q 4-hydroxy- HOGA1
9,13304 0 Q86YB7;Q Q86YB7;Q Enoyl-CoA ECHDC2
2,47597 0,011646 Q8IUX7;Q Q8IUX7;Q Adipocyte AEBP1
3,27991 0,004072 Q8IXI1;L Q8IXI1 Mitochond RHOT2
1,30017 0,072368 Q8IYB8;B1 Q8IYB8 ATP-depen SUPV3L1
1,43999 0,057348 Q8IYS1 Q8IYS1 Peptidase I PM20D2
3,12737 0,004898 Q8IZ83-3;C Q8IZ83-3;C Aldehyde c ALDH16A1
2,31713 0,014407 Q8N0W3;C Q8N0W3;C L-fucose ki FUK
5,0356 0,000517 Q8N0X4;Q Q8N0X4;Q Citrate lya CLYBL
12,0892 0 Q8N142;Q Q8N142;Q Adenylosu ADSSL1
3,58769 0,002786 Q8N163-2; Q8N163-2; Cell cycle a CCAR2
8,88232 0 Q8N335;C Q8N335;C Glycerol-3- GPD1L
2,32782 0,014187 Q8N4T8;Q Q8N4T8;Q Carbonyl r CBR4
3,61685 0,002694 Q8N8N7;G Q8N8N7;G Prostaglan PTGR2
14,9079 0 Q8N8Y2;E Q8N8Y2 V-type pro ATP6V0D2
3,2683 0,004119 Q8NBN7;C Q8NBN7;C Retinol del RDH13
7,14776 4,3E-05 Q8NBS9;Q Q8NBS9;Q Thioredoxi TXNDC5
4,34661 0,001094 Q8NBU5;C Q8NBU5;C ATPase fan ATAD1
6,49659 0,000105 Q8NBX0 Q8NBX0 Saccharopi SCCPDH
2,75177 0,007988 Q8NCN5;A Q8NCN5;A Pyruvate d PDPR
1,92543 0,026245 Q8NCW5-2 Q8NCW5-2 NAD(P)H-h APOA1BP
7,51604 2,8E-05 Q8NDH3;C Q8NDH3;C Probable a NPEPL1
7,24245 3,52E-05 Q8NE62;C Q8NE62 Choline de CHDH
2,07639 0,020859 Q8NE86-3; Q8NE86-3; Calcium un MCU
9,14524 0 Q8NFV4;Q Q8NFV4;Q Alpha/bet ABHD11
4,35858 0,00108 Q8NFW8;C Q8NFW8;C N-acylneur CMAS
3,44119 0,003405 Q8TC12-2; Q8TC12-2; Retinol del RDH11
5,55092 0,000376 Q8TCS8;H Q8TCS8 Polyribonu PNPT1
2,15581 0,018325 Q8WUM4; Q8WUM4; Programm PDCD6IP
6,47081 0,000104 Q8WW52; Q8WW52; Protein FAI FAM151A
2,60551 0,009662 Q8WX93-3 Q8WX93-3 Palladin PALLD
0,60821 0,248539 Q92499;F1 Q92499;F1 ATP-depen DDX1
7,42813 2,75E-05 Q92506 Q92506 Estradiol 1 HSD17B8
1,68869 0,038144 Q92523-3; Q92523-3; Carnitine C CPT1B
11,1102 0 Q92597;Q Q92597;Q Protein ND NDRG1
2,31314 0,014485 Q92616 Q92616 Translatior GCN1L1
0,705253 0,20653 Q92626;H Q92626 Peroxidasi PDXN
6,65771 7,06E-05 Q92820 Q92820 Gamma-gli GGH
6,37736 0,000139 Q92896;Q Q92896;Q Golgi appa GLG1
6,48271 0,000104 Q92947;Q Q92947;Q Glutaryl-Cc GCDH
2,17995 0,017604 Q93009-3; Q93009-3; Ubiquitin c USP7
8,0023 1,03E-05 Q93050-1; Q93050-1; V-type pro ATP6V0A1
19,6602 0 Q93088;E5 Q93088;E5 Betaine--h BHMT
6,04136 0,000193 Q93099;H Q93099 Homogent HGD
2,7929 0,007593 Q969I3-2;C Q969I3-2;C Glycine N- GLYATL1
7,61176 1,9E-05 Q969Z3;Q Q969Z3;Q Mitochond MARC2
2,79338 0,007593 Q96AB3;K Q96AB3;K Isochorism ISOC2
7,10511 4,26E-05 Q96AC1;Q Q96AC1;Q Fermitin fa FERMT2
10,6405 0 Q96BM9;A Q96BM9 ADP-ribosy ARL8A
1,2021 0,085383 Q96BW5;C Q96BW5;C Phosphotri PTER

7,35326 2,71E-05 Q96C23;H: Q96C23;H: Aldose 1-e GALM
13,9349 0 Q96CM8-3 Q96CM8-3 Acyl-CoA s' ACSF2
3,24378 0,004231 Q96CX2;Q: Q96CX2 BTB/POZ d KCTD12
3,42964 0,003434 Q96DB5;H: Q96DB5;H: Regulator (RMDN1
10,6424 0 Q96DC8;Q Q96DC8;Q Enoyl-CoA ECHDC3
20,1807 0 Q96DG6 Q96DG6 Carboxymε CMBL
3,13472 0,004825 Q96DV4;Q Q96DV4 39S riboso MRPL38
0,997204 0,121919 Q96EK6;G: Q96EK6;G: Glucosami GNPAT1
7,74991 1,98E-05 Q96EY1-2; Q96EY1-2; Dnal homc DNAJA3
2,27649 0,015266 Q96EY7;Q: Q96EY7 Pentatricol PTC3
0,59735 0,253237 Q96FJ2 Q96FJ2 Dynein lig† DYNLL2
3,80602 0,002134 Q96G03;E: Q96G03 Phosphogl† PGM2
4,53332 0,000863 Q96GK7;Q Q96GK7;Q Fumarylac† FAHD2A;FAHD2B
1,80464 0,0317 Q96HC4;Q Q96HC4;Q PDZ and LI PDLIM5
2,25822 0,01572 Q96HE7;G: Q96HE7;G: ERO1-like † ERO1L
0,722154 0,200363 Q96HY7;H: Q96HY7 Probable 2 DHTKD1
6,0476 0,000194 Q96I99;Q9 Q96I99;Q9 Succinyl-Cc SUCLG2
0,601533 0,251333 Q96IU4;F8 Q96IU4;F8 Alpha/betε ABHD14B
11,5327 0 Q96IX5 Q96IX5 Up-regulat USMG5
9,65494 0 Q96KP4;Q: Q96KP4;Q: Cytosolic n CNDP2
3,87094 0,001959 Q96LJ7;H0 Q96LJ7;H0 Dehydroge DHRS1
9,60472 0 Q96NB2;R: Q96NB2 Sideroflexi SFXN2
0,006833 0,975406 Q96QK1;I3 Q96QK1 Vacuolar p VPS35
4,43687 0,000987 Q96RP9;Q: Q96RP9;Q: Elongation GFM1
0,906996 0,142531 Q99436;Q: Q99436;Q: Proteasom PSMB7
3,25394 0,004194 Q99460-2; Q99460-2; 26S protea PSMD1
0,683431 0,214932 Q99497;K7 Q99497;K7 Protein de† PARK7
9,54236 0 Q99536;Q: Q99536;Q: Synaptic ve† VAT1
6,56196 9,23E-05 Q99538-3; Q99538-3; Legumain LGMN
14,6588 0 Q99541;Q: Q99541 Perilipin-2 PLIN2
9,92405 0 Q99714;Q: Q99714;Q: 3-hydroxyε HSD17B10
1,91479 0,026605 Q99733;Q: Q99733;Q: Nucleosom NAP1L4
1,45118 0,056367 Q99797 Q99797 Mitochond MIPEP
17,0196 0 Q99798;A: Q99798;A: Aconitate † ACO2
2,7368 0,008147 Q99832;Q: Q99832;Q: T-complex CCT7
4,39946 0,001052 Q9BPW8;† Q9BPW8;† Protein Ni† NIPSNAP1
1,66941 0,039338 Q9BPX5 Q9BPX5 Actin-relat ARPC5L
12,6649 0 Q9BQ69 Q9BQ69 O-acetyl-A MACROD1
5,54976 0,000375 Q9BQ95;J: Q9BQ95;J: Evolutiona ECSIT
2,6547 0,009052 Q9BQE3;F: Q9BQE3;F: Tubulin al† TUBA1C
1,78193 0,032736 Q9BR76;A: Q9BR76 Coronin-1E CORO1B
1,23171 0,081264 Q9BRA2;I3 Q9BRA2;I3 Thioredoxi TXNDC17
1,1487 0,093177 Q9BRF8;Q: Q9BRF8;Q: Serine/thrε CPPED1
2,4465 0,012158 Q9BS26 Q9BS26 Endoplasm ERP44
16,017 0 Q9BSE5 Q9BSE5 Agmatinas AGMAT
5,13925 0,000474 Q9BSH5 Q9BSH5 Haloacid d† HDHD3
6,5038 0,000105 Q9BSJ8-2;† Q9BSJ8-2;† Extended s ESYT1
14,2331 0 Q9BTV4;V: Q9BTV4 Transmem TMEM43
3,97826 0,001664 Q9BTZ2;Q: Q9BTZ2;Q: Dehydroge DHRS4;DHRS4L2
5,19531 0,00045 Q9BUF5;K: Q9BUF5;K: Tubulin be† TUBB6

7,14254 4,29E-05 Q9BUT1;Q Q9BUT1;Q 3-hydroxyl BDH2
8,49284 0 Q9BV79;Q Q9BV79;Q Trans-2-en MECR
1,59809 0,044382 Q9BVA1 Q9BVA1 Tubulin be TUBB2B
1,53648 0,048994 Q9BVK6 Q9BVK6 Transmem TMED9
0,537784 0,285676 Q9BX66-1; Q9BX66-1; Sorbin and SORBS1
5,45358 0,000406 Q9BX68 Q9BX68 Histidine tr HINT2
2,70212 0,008528 Q9BXJ7;HC Q9BXJ7;HC Protein am AMN
0,478713 0,319644 Q9B XK5-2; Q9B XK5-2; Bcl-2-like p BCL2L13
4,14672 0,001389 Q9BXN1;Q Q9BXN1;Q Asporin ASPN
1,5387 0,048894 Q9BY32-2; Q9BY32-2; Inosine triph ITPA
1,96737 0,02462 Q9BY49;B Q9BY49;B Peroxisom PECP
2,48342 0,011513 Q9BYD1;E Q9BYD1;E 39S riboso MRPL13
4,57539 0,000833 Q9BYD6;H Q9BYD6;H 39S riboso MRPL1
4,60621 0,000793 Q9BYV1;Q Q9BYV1;Q Alanine--gl AGXT2
2,99332 0,005846 S4R369;Q S4R369;Q 39S riboso MRPL37
2,35564 0,013765 Q9BZK7;A Q9BZK7;A F-box-like/ TBL1XR1
9,46301 0 Q9GZM7;C Q9GZM7;C Tubulointe TINAGL1
7,52458 2,81E-05 Q9H008;Q Q9H008 Phospholy: LHPP
3,27901 0,004072 Q9H061;Q Q9H061;Q Transmem TMEM126A
4,4705 0,000928 Q9H0E2;E Q9H0E2;E Toll-intera TOLLIP
3,54621 0,002927 Q9H0U4;E Q9H0U4;E Ras-relate RAB1B
8,26285 0 Q9H0W9;C Q9H0W9;C Ester hydr C11orf54
7,77599 1,99E-05 Q9H223 Q9H223 EH domain EHD4
10,2434 0 Q9H299;Q Q9H299;Q SH3 domai SH3BGR13
3,42896 0,003435 Q9H2A2-2; Q9H2A2-2; Aldehyde c ALDH8A1
1,78127 0,032742 Q9H2M3;C Q9H2M3;C S-methylm BHMT2
8,53906 0 Q9H2U2;Q Q9H2U2;Q Inorganic p PPA2
2,02788 0,022476 Q9H2W6;A Q9H2W6 39S riboso MRPL46
3,44906 0,003355 Q9H3G5;H Q9H3G5;H Probable s CPVL
1,71006 0,036903 Q9H3K2 Q9H3K2 Growth ho GHITM
3,97819 0,001662 Q9H479 Q9H479 Fructosam FN3K
6,0876 0,000182 Q9H4A4;A Q9H4A4;A Aminopept RNPEP
4,19727 0,001279 Q9H4M9;A Q9H4M9;A EH domain EHD1
7,86171 2,02E-05 Q9H7Z7;A Q9H7Z7;A Prostaglan PTGES2
3,99897 0,001629 Q9H845;H Q9H845;H Acyl-CoA d ACAD9
4,09239 0,001474 Q9H8L6;V Q9H8L6 Multimerir MMRN2
5,28827 0,000446 Q9H9B4;D Q9H9B4;D Sideroflexi SFXN1
1,61827 0,042939 Q9H9P8;C Q9H9P8;C L-2-hydrox L2HGDH
3,72263 0,002371 Q9HA77;H Q9HA77 Probable c CARS2
6,1114 0,000183 Q9HAT2;Q Q9HAT2;Q Sialate O-a SIAE
2,31137 0,014508 Q9HAV0;C Q9HAV0 Guanine nt GNB4
14,3818 0 Q9HBG4;A Q9HBG4 V-type pro ATP6V0A4
2,87652 0,006854 Q9HBL7 Q9HBL7 Plasminog PLGRKT
10,4241 0 Q9HCC0;Q Q9HCC0;Q Methylcroi MCCC2
2,40792 0,012867 Q9HDC9;H Q9HDC9;H Adipocyte APMAP
2,54168 0,01064 Q9NP92 Q9NP92 28S riboso MRPS30
1,86626 0,028675 Q9NPH2-2 Q9NPH2-2 Inositol-3- I SYNA1
5,51485 0,000372 Q9NPJ3-2; Q9NPJ3-2; Acyl-coenz ACOT13
1,44469 0,056898 Q9NQC3-3 Q9NQC3-3 Reticulon- RTN4
9,90328 0 Q9NQR4;H Q9NQR4;H Omega-am NIT2

3,98375 0,001657 Q9NQT8;E Q9NQT8;E Kinesin-like KIF13B
3,87889 0,001954 Q9NQX3;C Q9NQX3;C Gephyrin;M GPHN
2,44736 0,012147 Q9NR12;Q Q9NR12;Q PDZ and LIM PDLIM7
4,63542 0,000786 Q9NR19;Q Q9NR19;Q Acetyl-coe ACSS2
2,4056 0,012869 Q9NR28-2; Q9NR28-2; Diablo homolog DIABLO
3,18694 0,00453 Q9NR45;Q Q9NR45 Sialic acid s NANS
5,16704 0,000459 Q9NRG7;G Q9NRG7 Epimerase SDR39U1
1,32713 0,069461 Q9NRV9;A Q9NRV9;A Heme-binding HEBP1
2,36766 0,013499 Q9NRW3 Q9NRW3 DNA dC->dAPOBEC3C
6,51346 0,000106 Q9NRX4;Q Q9NRX4;Q 14 kDa phospho PHPT1
0,835228 0,16322 Q9NSD9;Q Q9NSD9;Q Phenylalanine FARSB
6,35451 0,000138 Q9NSE4 Q9NSE4 Isoleucine- IARS2
6,46512 0,000104 Q9NTJ5;E9 Q9NTJ5;E9 Phosphatidyl SACM1L
7,82711 2,01E-05 Q9NUB1;C Q9NUB1;C Acetyl-coe ACSS1
12,8123 0 Q9NUJ1;Q Q9NUJ1;Q Mycophenolate ABHD10
3,13678 0,004831 Q9NVI7-2; Q9NVI7-2; ATPase family ATAD3A
4,9715 0,000533 Q9NVJ2;Q Q9NVJ2 ADP-ribosyl ARL8B
2,41137 0,012825 Q9NWU1; C Q9NWU1; C 3-oxoacyl- OXSM
9,21666 0 Q9NX40;Q Q9NX40;Q OCIA domain OCIAD1
1,65633 0,040199 Q9NX46 Q9NX46 Poly(ADP-ribose) ADPRHL2
5,91734 0,000243 Q9NXA8-2 Q9NXA8-2 NAD-dependent SIRT5
3,8929 0,001909 Q9NYL9;H Q9NYL9;H Tropomodulin TMOD3
13,901 0 Q9NYQ3-2 Q9NYQ3-2 Hydroxyacyl HAO2
2,57089 0,010149 Q9NYU2;Q Q9NYU2;Q UDP-glucosyl UGGT1
1,87552 0,028346 Q9NZ08-2; Q9NZ08-2; Endoplasmic ERAP1
11,2419 0 Q9NZ45 Q9NZ45 CDGSH iron CISD1
3,10141 0,005061 Q9NZB2-4; Q9NZB2-4; Constitutive FAM120A
2,69279 0,008647 Q9NZL9;Q Q9NZL9;Q Methionine MAT2B
9,98486 0 Q9NZM1-6 Q9NZM1-6 Myoferlin MYOF
12,5557 0 Q9NZN4;Q Q9NZN4;Q EH domain EHD2
3,87324 0,001962 Q9NZU0 Q9NZU0 Leucine-rich FLRT3
4,12106 0,001419 Q9P015;E5 Q9P015;E5 39S ribosomal MRPL15
3,06826 0,005331 Q9P0J0-2; C Q9P0J0-2; C NADH dehydrogenase NDUFA13
2,75475 0,007949 Q9P0J1;Q Q9P0J1;Q [Pyruvate] PDP1
6,7801 7,23E-05 Q9P0L0;Q Q9P0L0;Q Vesicle-associated VAPA
2,52847 0,010819 Q9P0M6;C Q9P0M6;C Core histone H2AFY2
10,8855 0 Q9P0Z9;K7 Q9P0Z9 Peroxisomal PIPOX
17,9442 0 Q9UBD6;A Q9UBD6;A Ammonium RHCG
3,67372 0,002518 Q9UBF2;Q Q9UBF2;Q Coatamer COPG2
6,28769 0,000158 Q9UBI6 Q9UBI6 Guanine nucleotide GNG12
7,31898 3,6E-05 Q9UBQ7;U Q9UBQ7;U Glyoxylate GRHPR
9,98002 0 Q9UBR2 Q9UBR2 Cathepsin B CTSZ
2,55343 0,010435 Q9UDR5;F Q9UDR5;F Alpha-amylase AASS
3,12804 0,004899 Q9UDW1; C Q9UDW1; C Cytochrome UQCR10
0,758629 0,187258 Q9UEY8-2; Q9UEY8-2; Gamma-aminobutyrate ADD3
13,4473 0 Q9UF12;K7 Q9UF12 Probable pyruvate PRODHD2
12,8573 0 Q9UFN0;Q Q9UFN0 Protein Nip3 NIPSNAP3A
3,45381 0,003319 Q9UGI8-2; Q9UGI8-2; Testin TES
2,98035 0,005933 Q9UGT4 Q9UGT4 Sushi domain SUSD2
3,10568 0,005038 Q9UH99-3 Q9UH99-3 SUN domain SUN2

0,825503 0,165996 Q9UHD8;C Q9UHD8;C Septin-9 SEPT9
6,76795 7,21E-05 Q9UHE5;Q Q9UHE5 N-acetyltra NAT8
3,47789 0,003221 Q9UHG3;C Q9UHG3;C Prenylcyst PCYOX1
7,10089 4,26E-05 Q9UHI7;Q Q9UHI7;Q Solute carr SLC23A1
14,576 0 Q9UHK6;Q Q9UHK6;Q Alpha-met AMACR
1,94445 0,02553 Q9UHL4;R Q9UHL4 Dipeptidyl DPP7
13,262 0 Q9UHQ9;F Q9UHQ9;F NADH-cytc CYB5R1
6,7856 7,27E-05 Q9UHY7;A Q9UHY7;A Enolase-ph ENOPH1
8,56158 0 Q9UI09;F8 Q9UI09 NADH deh' NDUFA12
18,7102 0 Q9UI12-2;I Q9UI12-2;I V-type pro ATP6V1H
9,94444 0 Q9UI17;Q Q9UI17;Q Dimethylgl DMGDH
1,94046 0,025666 Q9UII2 Q9UII2 ATPase inh ATPIF1
14,157 0 Q9UIJ7;Q9 Q9UIJ7;Q9 GTP:AMP f AK3
6,50307 0,000105 Q9UJ70;Q Q9UJ70;Q N-acetyl-D NAGK
6,12681 0,000176 Q9UJS0-2;I Q9UJS0-2;I Calcium-bi SLC25A13
1,00892 0,119758 Q9UJU6;F Q9UJU6;F Drebrin-lik DBNL
17,1081 0 Q9UJW2;C Q9UJW2;C Tubulointe TINAG
12,6818 0 Q9UJZ1;Q Q9UJZ1;Q Stomatin-li STOML2
4,62436 0,000796 Q9UKK3 Q9UKK3 Poly [ADP- PARP4
6,81 7,29E-05 Q9UKU7;Q Q9UKU7;Q Isobutyryl- ACAD8
12,6304 0 Q9UL12-2; Q9UL12-2; Sarcosine c SARDH
1,08256 0,104761 Q9UL25 Q9UL25 Ras-relatec RAB21
16,9905 0 Q9ULD0;Q Q9ULD0;Q 2-oxogluta OGDHL
13,966 0 Q9ULV4;Q Q9ULV4;Q Coronin-1C CORO1C
2,0264 0,022494 Q9UMS4;F Q9UMS4;F Pre-mRNA PRPF19
7,04608 5,05E-05 Q9UN36;G Q9UN36;G Protein ND NDRG2
5,51053 0,000378 Q9UNF0-2 Q9UNF0-2 Protein kin PACSIN2
1,00309 0,120699 Q9UNZ2-6 Q9UNZ2-6 NSFL1 cofa NSFL1C
3,10448 0,005033 Q9UQ80;C Q9UQ80;C Proliferatic PA2G4
0,431561 0,351613 Q9UQE7;D Q9UQE7 Structural i SMC3
1,02538 0,116487 Q9Y230;Q Q9Y230;Q RuvB-like 2 RUVBL2
0,538273 0,285588 Q9Y265;Q Q9Y265;Q RuvB-like 1 RUVBL1
4,46828 0,000937 Q9Y276;C Q9Y276;C Mitochond BCS1L
5,4715 0,000394 Q9Y277;Q Q9Y277;Q Voltage-de VDAC3
2,9458 0,006217 Q9Y281;Q Q9Y281;Q Cofilin-2 CFL2
0,233418 0,53949 Q9Y2D0;C Q9Y2D0 Carbonic a CA5B
12,4635 0 Q9Y2Q3;Q Q9Y2Q3;Q Glutathion GSTK1
10,8517 0 Q9Y2R0;K Q9Y2R0;K Cytochrom COA3
3,5236 0,003036 Q9Y2S2-2;I Q9Y2S2-2;I Lambda-cr CRYL1
3,27351 0,004102 Q9Y2T3;Q Q9Y2T3;Q Guanine d GDA
1,73033 0,035756 Q9Y2X3;H Q9Y2X3;H Nucleolar f NOP58
1,28057 0,074813 Q9Y2Z0-2;I Q9Y2Z0-2;I Suppressor SUGT1
3,40302 0,003512 Q9Y305-2; Q9Y305-2; Acyl-coenz ACOT9
11,861 0 Q9Y394-2; Q9Y394-2; Dehydroge DHRS7
2,85414 0,007086 Q9Y3D6 Q9Y3D6 Mitochond FIS1
0,3856 0,387069 Q9Y3I0 Q9Y3I0 tRNA-splici RTCB
4,08789 0,001475 Q9Y3Z3;AC Q9Y3Z3;AC Deoxynucl SAMHD1
10,3764 0 Q9Y490 Q9Y490 Talin-1 TLN1
6,53086 0,000107 Q9Y4F1;C Q9Y4F1;C FERM, Rho FARP1
4,49393 0,000899 Q9Y4G6;A Q9Y4G6;A Talin-2 TLN2

11,8104	0	Q9Y4W6;K	Q9Y4W6	AFG3-like Ꞥ	AFG3L2
5,08355	0,00048	Q9Y512	Q9Y512	Sorting anc	SAMM50
12,4073	0	Q9Y5U8;Q	Q9Y5U8;Q	Mitochond	MPC1;BRP44L
2,36879	0,013482	Q9Y5X3;Q!	Q9Y5X3;Q!	Sorting ne	SNX5
11,6021	0	Q9Y617;Q!	Q9Y617;Q!	Phosphose	PSAT1
0,691725	0,211556	Q9Y619;F2	Q9Y619	Mitochond	SLC25A15
4,7283	0,000676	Q9Y678;H!	Q9Y678	Coatome	COPG1
19,6805	0	Q9Y696;A!	Q9Y696	Chloride in	CLIC4
5,50914	0,000378	Q9Y697;Q!	Q9Y697;Q!	Cysteine di	NFS1
0,433159	0,350687	Q9Y6C2;A!	Q9Y6C2	EMILIN-1	EMILIN1
7,69013	1,96E-05	Q9Y6C9;E!	Q9Y6C9;E!	Mitochond	MTCH2
5,17354	0,000448	Q9Y6N5;H!	Q9Y6N5;H!	Sulfide:qui	SQRDL
8,76737	0	Q9Y6R1-3;	Q9Y6R1-3;	Electrogen	SLC4A4
14,6245	0	Q9Y6U3;Q	Q9Y6U3;Q	Adseverin	SCIN
1,95712	0,025024	R4GNH3;P	R4GNH3;P	26S protea	PSMC3

2BF;HIST1H2BC;HIST1H2BD;H2BFS;HIST1H2BK;HIST1H2BL

