

**Table S1** - List of differentially abundant proteins in PDL of the control and obese groups. Average peak area values of each group, p-values (*t*-test), and fold-change (obese / control) are shown.

Accession	Description	Protein Symbol	Peak area										t-test	Fold-Change
			Control 1	Control 2	Control 3	Control Average	Obese 1	Obese 2	Obese 3	Obese 4	Obese 5	Obese Average		
Up-regulated in Obesity														
Q9EQP5	Prolargin	Prelp	0.0E0	0.0E0	0.0E0	0.0E0	5.4E6	1.3E7	1.8E7	1.5E7	6.3E6	1.2E7	0.014	1.2E7
Q4KLF8	Actin-related protein 2/3 complex subunit 5	Arcp5	0.0E0	0.0E0	0.0E0	0.0E0	1.0E7	9.1E6	0.0E0	7.4E6	5.4E6	6.4E6	0.036	6.4E6
I6L9G6	RCG31562, isoform CRA_c	I6l9g6	0.0E0	0.0E0	0.0E0	0.0E0	4.0E6	3.6E6	1.7E6	2.4E6	0.0E0	2.4E6	0.049	2.4E6
Q9EPC6	Profilin-2	Prof2	0.0E0	0.0E0	0.0E0	0.0E0	3.1E6	3.0E6	3.4E6	0.0E0	1.3E6	2.2E6	0.046	2.2E6
A3FM27	Chloride intracellular channel protein	A3fm27	0.0E0	0.0E0	0.0E0	0.0E0	1.4E6	1.5E6	1.4E6	2.6E6	0.0E0	1.4E6	0.043	1.4E6
Q5XFW8	Protein SEC13 homolog	Sec13	0.0E0	2.2E6	0.0E0	7.4E5	2.7E6	5.9E6	5.0E6	6.2E6	3.0E6	4.6E6	0.014	6.1
O35796	Complement component 1 Q subcomponent-binding protein, mitochondrial	C1qbp	0.0E0	2.9E6	0.0E0	9.5E5	3.4E6	4.0E6	2.5E6	6.1E6	4.7E6	4.1E6	0.026	4.3
P07895	Superoxide dismutase	Sod2	0.0E0	3.5E6	4.3E6	2.6E6	4.9E6	4.7E6	1.1E7	8.6E6	9.4E6	7.8E6	0.040	3.0
P17764	Acetyl-CoA acetyltransferase, mitochondrial	Thil	0.0E0	1.6E6	3.1E6	1.6E6	4.0E6	4.8E6	3.8E6	6.1E6	4.5E6	4.6E6	0.011	3.0
P16975	SPARC	Sprc	1.1E7	1.5E7	8.2E6	1.1E7	1.5E7	2.6E7	1.8E7	1.9E7	1.7E7	1.9E7	0.042	1.7
Down-regulated in Obesity														
A0A0G2K0W9	Proteasome subunit alpha type	A0a0g2k0w9	4.4E6	4.8E6	3.4E6	4.2E6	3.3E6	2.9E6	3.2E6	3.6E6	2.2E6	3.0E6	0.037	-1.4
A0A0U1RRV7	RCG61099, isoform CRA_b	A0a0u1rrv7	1.3E7	1.3E7	1.6E7	1.4E7	8.5E6	8.1E6	1.2E7	1.1E7	8.0E6	9.5E6	0.018	-1.5
M0RA26	Ribosomal protein S27-like	M0ra26	8.3E6	1.1E7	8.2E6	9.2E6	8.2E6	7.0E6	5.1E6	6.4E6	4.2E6	6.2E6	0.044	-1.5
D3ZWE0	Histone H2A	D3zwe0	6.5E8	6.5E8	6.1E8	6.4E8	4.5E8	4.1E8	4.9E8	5.4E8	2.4E8	4.3E8	0.022	-1.5
P61805	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit DAD1	Dad1	9.1E6	7.9E6	5.4E6	7.5E6	5.2E6	4.6E6	6.1E6	5.0E6	3.3E6	4.9E6	0.039	-1.5
A0A0G2K757	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2	A0a0g2k757	2.7E7	2.4E7	1.7E7	2.2E7	1.5E7	1.5E7	1.3E7	2.0E7	9.1E6	1.4E7	0.049	-1.5

Q794E4	Heterogeneous nuclear ribonucleoprotein F	Hnrpf	1.8E7	1.9E7	1.5E7	<b>1.7E7</b>	1.4E7	9.5E6	1.1E7	9.9E6	1.1E7	<b>1.1E7</b>	0.002	<b>-1.6</b>
Q68FP1	Gelsolin	Gels	6.9E7	6.7E7	4.3E7	<b>6.0E7</b>	4.2E7	4.4E7	3.3E7	4.2E7	2.5E7	<b>3.7E7</b>	0.028	<b>-1.6</b>
F1M779	Clathrin heavy chain	F1m779	4.8E7	4.5E7	2.7E7	<b>4.0E7</b>	2.6E7	2.7E7	1.9E7	3.2E7	2.0E7	<b>2.5E7</b>	0.041	<b>-1.6</b>
Q4V7D1	Signal sequence receptor, alpha	Q4v7d1	2.1E7	2.5E7	1.7E7	<b>2.1E7</b>	1.8E7	1.1E7	1.2E7	1.4E7	7.8E6	<b>1.3E7</b>	0.026	<b>-1.7</b>
Q6AXS5	Plasminogen activator inhibitor 1 RNA-binding protein	Pairb	3.8E6	4.9E6	3.6E6	<b>4.1E6</b>	2.5E6	2.4E6	2.1E6	3.9E6	1.3E6	<b>2.4E6</b>	0.038	<b>-1.7</b>
P62909	40S ribosomal protein S3	Rs3	5.0E7	4.3E7	3.4E7	<b>4.2E7</b>	2.8E7	2.8E7	2.4E7	2.7E7	1.5E7	<b>2.4E7</b>	0.009	<b>-1.7</b>
P35427	60S ribosomal protein L13a	RI13a	1.9E7	2.3E7	1.4E7	<b>1.9E7</b>	1.1E7	1.2E7	9.5E6	1.3E7	7.3E6	<b>1.1E7</b>	0.019	<b>-1.7</b>
F1LSW7	60S ribosomal protein L14	F1lsw7	2.9E7	2.8E7	1.9E7	<b>2.5E7</b>	1.7E7	1.4E7	1.3E7	2.0E7	7.7E6	<b>1.4E7</b>	0.025	<b>-1.8</b>
P31000	Vimentin	Vime	5.2E8	6.5E8	4.3E8	<b>5.3E8</b>	2.8E8	2.8E8	2.6E8	4.6E8	2.3E8	<b>3.0E8</b>	0.019	<b>-1.8</b>
A0A0G2JUA5	AHNAK nucleoprotein	A0a0g2jua5	2.0E8	2.0E8	9.5E7	<b>1.6E8</b>	1.1E8	8.1E7	8.6E7	1.2E8	6.7E7	<b>9.3E7</b>	0.047	<b>-1.8</b>
Q6AY09	Heterogeneous nuclear ribonucleoprotein H2	Hnrh2	2.1E7	2.7E7	2.0E7	<b>2.3E7</b>	1.4E7	1.5E7	1.7E7	1.1E7	6.9E6	<b>1.3E7</b>	0.013	<b>-1.8</b>
Q63030	Rat alpha-smooth muscle actin mRNA (Fragment)	Q63030	1.6E9	2.0E9	1.0E9	<b>1.5E9</b>	1.1E9	8.2E8	7.5E8	9.9E8	5.4E8	<b>8.4E8</b>	0.030	<b>-1.8</b>
Q5PQK2	FUS RNA-binding protein	Q5pqk2	6.9E6	8.2E6	5.0E6	<b>6.7E6</b>	4.2E6	3.3E6	3.9E6	4.2E6	2.9E6	<b>3.7E6</b>	0.008	<b>-1.8</b>
Q66H80	Coatomer subunit delta	Copd	8.2E6	8.1E6	4.7E6	<b>7.0E6</b>	3.6E6	4.3E6	3.5E6	5.3E6	2.6E6	<b>3.9E6</b>	0.022	<b>-1.8</b>
Q6PDU1	Serine/arginine-rich splicing factor 2	Srsf2	1.6E7	1.5E7	1.2E7	<b>1.4E7</b>	8.3E6	8.3E6	6.2E6	1.1E7	4.9E6	<b>7.8E6</b>	0.009	<b>-1.8</b>
A0A0G2JSH9	Peroxiredoxin-2	A0a0g2jsh9	3.3E7	3.5E7	1.8E7	<b>2.9E7</b>	1.6E7	1.2E7	1.7E7	2.4E7	9.7E6	<b>1.6E7</b>	0.040	<b>-1.8</b>
Q9EPB1	Dipeptidyl peptidase 2	Dpp2	3.4E7	3.1E7	1.8E7	<b>2.8E7</b>	1.8E7	2.1E7	1.4E7	1.7E7	5.0E6	<b>1.5E7</b>	0.048	<b>-1.8</b>
A0A0G2JTG7	Heterogeneous nuclear ribonucleoprotein H	A0a0g2jtg7	2.4E7	3.1E7	1.7E7	<b>2.4E7</b>	1.5E7	1.3E7	1.7E7	1.2E7	6.9E6	<b>1.3E7</b>	0.027	<b>-1.8</b>
F1LNF1	Heterogeneous nuclear ribonucleoproteins A2/B1	F1lnf1	8.0E7	7.7E7	5.3E7	<b>7.0E7</b>	3.3E7	3.6E7	3.6E7	6.4E7	2.0E7	<b>3.8E7</b>	0.029	<b>-1.9</b>
P29975	Aquaporin-1	Aqp1	1.6E7	2.2E7	9.9E6	<b>1.6E7</b>	7.5E6	7.9E6	9.5E6	1.2E7	5.2E6	<b>8.5E6</b>	0.049	<b>-1.9</b>
F1M7X3	Cadherin 13	F1m7x3	1.2E7	1.2E7	9.5E6	<b>1.1E7</b>	8.1E6	4.2E6	6.7E6	5.7E6	4.5E6	<b>5.9E6</b>	0.004	<b>-1.9</b>
A0A0H2UHT6	Ribosomal protein S18	A0a0h2uht6	4.2E7	3.9E7	2.7E7	<b>3.6E7</b>	1.7E7	2.0E7	1.4E7	3.0E7	1.3E7	<b>1.9E7</b>	0.018	<b>-1.9</b>
Q3KRF2	High density lipoprotein binding protein (Vigilin)	Q3krf2	1.8E7	1.8E7	1.1E7	<b>1.6E7</b>	7.1E6	8.2E6	7.1E6	1.3E7	5.4E6	<b>8.2E6</b>	0.026	<b>-1.9</b>

Q6S3A0	Plectin	Q6s3a0	5.1E7	5.0E7	3.0E7	<b>4.4E7</b>	2.8E7	2.2E7	1.7E7	3.0E7	1.6E7	<b>2.3E7</b>	0.016	<b>-1.9</b>
P30427	Plectin	Plec	5.1E7	5.0E7	3.0E7	<b>4.4E7</b>	2.8E7	2.2E7	1.7E7	3.0E7	1.6E7	<b>2.3E7</b>	0.016	<b>-1.9</b>
Q68FR9	Elongation factor 1-delta	Ef1d	1.6E7	2.4E7	1.8E7	<b>1.9E7</b>	1.5E7	8.3E6	8.6E6	1.3E7	5.9E6	<b>1.0E7</b>	0.014	<b>-1.9</b>
P62250	40S ribosomal protein S16	Rs16	2.9E7	2.7E7	1.2E7	<b>2.3E7</b>	1.5E7	1.2E7	1.2E7	1.2E7	6.5E6	<b>1.2E7</b>	0.049	<b>-1.9</b>
Q0QEW8	Ribosomal protein L18 (Fragment)	Q0qew8	4.2E7	3.8E7	1.9E7	<b>3.3E7</b>	1.7E7	2.1E7	1.8E7	2.1E7	8.9E6	<b>1.7E7</b>	0.038	<b>-1.9</b>
A0A0G2KA35	Uncharacterized protein	A0a0g2ka35	7.2E7	7.6E7	4.9E7	<b>6.6E7</b>	2.8E7	3.1E7	3.8E7	6.0E7	1.3E7	<b>3.4E7</b>	0.038	<b>-1.9</b>
D4A6W6	Similar to 60S ribosomal protein L8	D4a6w6	3.0E7	3.0E7	1.6E7	<b>2.6E7</b>	1.3E7	1.3E7	1.1E7	2.3E7	5.9E6	<b>1.3E7</b>	0.044	<b>-2.0</b>
P05964	Protein S100-A6	S10a6	1.4E8	1.6E8	8.3E7	<b>1.3E8</b>	6.7E7	6.0E7	6.2E7	9.0E7	4.6E7	<b>6.5E7</b>	0.020	<b>-2.0</b>
G3V8L3	Lamin A, isoform CRA_b	G3v8l3	3.6E8	4.1E8	2.1E8	<b>3.3E8</b>	1.5E8	1.5E8	1.6E8	2.5E8	1.2E8	<b>1.7E8</b>	0.024	<b>-2.0</b>
P63018	Heat shock cognate 71 kDa protein	Hsp7c	1.4E8	1.6E8	6.8E7	<b>1.2E8</b>	4.9E7	5.4E7	6.5E7	9.2E7	3.9E7	<b>6.0E7</b>	0.040	<b>-2.0</b>
A0A0G2JY31	Alpha-1-antiproteinase	A0a0g2jy31	6.5E7	5.6E7	3.1E7	<b>5.1E7</b>	2.8E7	2.2E7	2.6E7	4.1E7	9.7E6	<b>2.5E7</b>	0.046	<b>-2.0</b>
B2RZ72	Actin-related protein 2/3 complex subunit 4	B2rz72	2.0E7	1.7E7	1.0E7	<b>1.6E7</b>	7.6E6	8.3E6	7.5E6	1.0E7	5.0E6	<b>7.7E6</b>	0.016	<b>-2.0</b>
Q66X93	Staphylococcal nuclease domain-containing protein 1	Snd1	2.3E7	1.9E7	1.4E7	<b>1.9E7</b>	9.3E6	1.1E7	7.4E6	1.4E7	3.4E6	<b>9.2E6</b>	0.025	<b>-2.0</b>
A0A0G2JZH3	Thrombospondin 3	A0a0g2jzh3	5.8E7	5.2E7	6.4E7	<b>5.8E7</b>	2.3E7	3.4E7	2.0E7	4.7E7	1.7E7	<b>2.8E7</b>	0.009	<b>-2.1</b>
A0A0G2JWX4	Keratin, type II cytoskeletal 2 epidermal	A0a0g2jwx4	3.5E7	2.1E7	1.9E7	<b>2.5E7</b>	1.8E7	1.7E7	1.3E7	3.9E6	9.5E6	<b>1.2E7</b>	0.045	<b>-2.1</b>
Q6PDV8	RCG31311	Q6pdv8	3.0E7	4.1E7	1.6E7	<b>2.9E7</b>	1.6E7	1.0E7	1.7E7	1.7E7	8.6E6	<b>1.4E7</b>	0.042	<b>-2.1</b>
I6L9G5	Rcn3 protein	I6l9g5	1.9E7	2.3E7	1.2E7	<b>1.8E7</b>	9.2E6	6.9E6	7.5E6	1.6E7	3.8E6	<b>8.6E6</b>	0.032	<b>-2.1</b>
Q05175	Brain acid soluble protein 1	Basp1	5.0E6	7.5E6	3.9E6	<b>5.5E6</b>	2.3E6	2.1E6	4.3E6	2.9E6	1.2E6	<b>2.6E6</b>	0.031	<b>-2.1</b>
F1LW91	Nuclear mitotic apparatus protein 1	F1lw91	3.1E6	2.4E6	2.5E6	<b>2.7E6</b>	9.8E5	1.1E6	2.5E6	1.4E6	1.9E5	<b>1.2E6</b>	0.035	<b>-2.1</b>
M0RA79	Uncharacterized LOC691828	M0ra79	1.0E8	9.3E7	1.0E8	<b>1.0E8</b>	4.7E7	3.4E7	5.7E7	6.7E7	2.8E7	<b>4.7E7</b>	0.002	<b>-2.1</b>
P08289	Alkaline phosphatase, tissue-nonspecific isozyme	Ppbt	4.7E7	5.1E7	2.8E7	<b>4.2E7</b>	2.2E7	1.6E7	2.9E7	1.9E7	1.1E7	<b>1.9E7</b>	0.012	<b>-2.2</b>
D3Z9E1	Elastin microfibril interfacier 1	D3z9e1	1.7E7	1.7E7	8.0E6	<b>1.4E7</b>	3.9E6	7.3E6	8.4E6	7.4E6	4.8E6	<b>6.3E6</b>	0.022	<b>-2.2</b>
P85845	Fascin	Fscn1	1.2E7	1.1E7	9.5E6	<b>1.1E7</b>	6.7E6	7.0E6	5.2E6	5.1E6	0.0E0	<b>4.8E6</b>	0.015	<b>-2.2</b>
A0A0G2JST3	Keratin, type II cytoskeletal 1	A0a0g2jst3	1.2E8	1.4E8	6.8E7	<b>1.1E8</b>	1.3E7	5.0E7	8.8E7	4.2E7	5.2E7	<b>4.9E7</b>	0.036	<b>-2.2</b>
G3V7U4	Lamin-B1	G3v7u4	7.8E7	6.9E7	4.3E7	<b>6.4E7</b>	2.1E7	2.6E7	2.9E7	4.9E7	1.7E7	<b>2.8E7</b>	0.016	<b>-2.2</b>

D3ZK97	Histone H3	D3zk97	2.1E8	2.3E8	4.2E8	<b>2.9E8</b>	1.6E8	2.0E8	1.2E8	1.2E8	2.2E7	<b>1.2E8</b>	0.046	<b>-2.3</b>
A0A0G2K9Q9	Keratin, type I cytoskeletal 17	A0a0g2k9q9	3.2E7	3.6E7	1.5E7	<b>2.8E7</b>	1.4E7	8.7E6	1.6E7	1.7E7	4.3E6	<b>1.2E7</b>	0.036	<b>-2.3</b>
A0A0G2KAM4	Fascin	A0a0g2kam4	1.2E7	1.1E7	9.5E6	<b>1.1E7</b>	6.6E6	7.0E6	4.3E6	5.1E6	0.0E0	<b>4.6E6</b>	0.012	<b>-2.3</b>
Q6LED0	Histone H3.1	H31	2.1E8	2.0E8	4.2E8	<b>2.8E8</b>	1.5E8	1.9E8	1.2E8	1.2E8	2.2E7	<b>1.2E8</b>	0.049	<b>-2.3</b>
D3ZJ08	Histone H3	D3zj08	2.4E8	2.2E8	4.2E8	<b>3.0E8</b>	1.6E8	1.9E8	1.2E8	1.2E8	2.0E7	<b>1.2E8</b>	0.029	<b>-2.4</b>
F1M013	60S ribosomal protein L7a	F1m013	3.3E7	3.2E7	2.2E7	<b>2.9E7</b>	1.4E7	1.1E7	3.9E6	1.8E7	1.2E7	<b>1.2E7</b>	0.005	<b>-2.5</b>
D3ZKU5	Similar to ribosomal protein L31	D3zku5	2.7E7	1.8E7	1.5E7	<b>2.0E7</b>	6.4E6	1.1E7	1.0E7	5.1E6	6.4E6	<b>7.8E6</b>	0.007	<b>-2.6</b>
F1LV13	Heterogeneous nuclear ribonucleoprotein M	F1lv13	1.4E7	1.6E7	8.2E6	<b>1.3E7</b>	4.1E6	4.3E6	3.8E6	9.1E6	3.1E6	<b>4.9E6</b>	0.012	<b>-2.6</b>
Q6IFV4	Keratin, type I cytoskeletal 13	K1c13	3.6E7	3.9E7	2.1E7	<b>3.2E7</b>	1.3E7	8.5E6	1.8E7	1.7E7	4.9E6	<b>1.2E7</b>	0.010	<b>-2.6</b>
F1LQ14	60S ribosomal protein L34	F1lq14	1.6E7	1.9E7	1.3E7	<b>1.6E7</b>	5.5E6	5.5E6	6.0E6	1.3E7	0.0E0	<b>6.0E6</b>	0.017	<b>-2.6</b>
D3ZBN0	Histone H1.5	H15	4.2E7	4.5E7	3.2E7	<b>4.0E7</b>	9.3E6	1.4E7	1.3E7	2.7E7	7.3E6	<b>1.4E7</b>	0.004	<b>-2.8</b>
G3V9R8	Heterogeneous nuclear ribonucleoprotein C	Hnrpc	1.2E7	4.9E6	6.2E6	<b>7.7E6</b>	4.6E6	3.0E6	0.0E0	4.1E6	2.1E6	<b>2.7E6</b>	0.042	<b>-2.8</b>
P02793	Ferritin light chain 1	Fril1	7.7E7	7.6E7	3.1E7	<b>6.1E7</b>	3.6E7	2.7E7	1.1E7	2.3E7	8.9E6	<b>2.1E7</b>	0.020	<b>-2.9</b>
A0A0G2JZS9	Uncharacterized protein	A0a0g2jzs9	7.8E7	4.2E7	7.4E7	<b>6.5E7</b>	0.0E0	2.4E7	3.1E7	3.4E7	1.5E7	<b>2.1E7</b>	0.009	<b>-3.1</b>
O35244	Peroxiredoxin-6	Prdx6	1.1E7	1.2E7	7.0E6	<b>9.9E6</b>	6.3E6	0.0E0	4.8E6	4.2E6	0.0E0	<b>3.1E6</b>	0.015	<b>-3.2</b>
A0A096MJE0	Keratin 32 (Fragment)	A0a096mje0	8.1E7	7.5E7	3.2E7	<b>6.3E7</b>	2.0E7	1.7E7	4.2E7	1.7E7	0.0E0	<b>1.9E7</b>	0.022	<b>-3.3</b>
A0A0H2UHU0	40S ribosomal protein S25	A0a0h2uhu0	2.9E7	2.3E7	1.1E7	<b>2.1E7</b>	1.0E7	0.0E0	4.7E6	1.4E7	3.3E6	<b>6.4E6</b>	0.026	<b>-3.3</b>
Q6IFU9	Keratin 16	Q6ifu9	7.7E7	6.9E7	7.9E7	<b>7.5E7</b>	1.9E7	1.1E7	3.1E7	3.1E7	1.6E7	<b>2.2E7</b>	0.0001	<b>-3.5</b>
Q5RKG9	Eukaryotic translation initiation factor 4B	Q5rk9	7.8E5	1.4E6	8.1E5	<b>1.0E6</b>	3.5E5	5.2E5	0.0E0	5.4E5	0.0E0	<b>2.8E5</b>	0.018	<b>-3.6</b>
A0A0G2K531	Glutathione peroxidase	A0a0g2k531	2.7E7	1.2E7	1.4E7	<b>1.8E7</b>	1.1E7	8.3E6	0.0E0	0.0E0	4.8E6	<b>4.7E6</b>	0.027	<b>-3.7</b>
D3ZZN4	Uncharacterized protein	D3zzn4	1.3E7	1.4E7	9.4E6	<b>1.2E7</b>	5.7E6	0.0E0	8.4E6	0.0E0	1.8E6	<b>3.2E6</b>	0.010	<b>-3.9</b>
A0A0G2JSL0	Proteasome subunit beta type	A0a0g2jsl0	1.1E7	1.4E7	6.5E6	<b>1.0E7</b>	5.0E6	4.6E6	0.0E0	0.0E0	3.0E6	<b>2.5E6</b>	0.009	<b>-4.1</b>
P52296	Importin subunit beta-1	lmb1	1.2E7	1.0E7	6.3E6	<b>9.6E6</b>	5.3E6	0.0E0	0.0E0	2.6E6	3.2E6	<b>2.2E6</b>	0.007	<b>-4.3</b>
B5DF91	ELAV-like protein 1	Elav1	1.0E7	9.9E6	3.9E6	<b>7.9E6</b>	0.0E0	0.0E0	3.1E6	5.4E6	0.0E0	<b>1.7E6</b>	0.025	<b>-4.6</b>
D4ABT8	Heterogeneous nuclear ribonucleoprotein U-like 2	D4abt8	2.5E6	3.3E6	1.7E6	<b>2.5E6</b>	0.0E0	0.0E0	1.1E6	1.5E6	0.0E0	<b>5.3E5</b>	0.013	<b>-4.7</b>

D3Z9G3	Uncharacterized protein	D3z9g3	5.1E7	6.4E7	2.6E7	<b>4.7E7</b>	1.9E7	2.2E7	0.0E0	8.6E6	0.0E0	<b>9.9E6</b>	0.012	<b>-4.7</b>
G3V9T1	Cysteine-rich secretory protein LCCL domain-containing 2	G3v9t1	4.8E6	4.8E6	2.4E6	<b>4.0E6</b>	2.3E6	0.0E0	1.9E6	0.0E0	0.0E0	<b>8.5E5</b>	0.013	<b>-4.7</b>
P62083	40S ribosomal protein S7	Rs7	2.4E7	3.6E7	7.1E6	<b>2.2E7</b>	9.9E6	8.0E6	0.0E0	4.9E6	0.0E0	<b>4.6E6</b>	0.037	<b>-4.9</b>
D3ZN79	Similar to 60S ribosomal protein L35	D3zn79	3.7E7	5.0E7	2.3E7	<b>3.7E7</b>	1.5E7	1.8E7	0.0E0	0.0E0	2.9E6	<b>7.2E6</b>	0.009	<b>-5.1</b>
A0A0G2K9X1	Secreted phosphoprotein 24	A0a0g2k9x1	1.4E7	1.8E7	4.4E6	<b>1.2E7</b>	0.0E0	8.8E6	0.0E0	2.5E6	0.0E0	<b>2.3E6</b>	0.038	<b>-5.3</b>
B2RYQ2	Serine/threonine-protein phosphatase 2A activator	B2ryq2	2.9E6	4.5E6	3.0E6	<b>3.5E6</b>	0.0E0	0.0E0	0.0E0	3.0E6	0.0E0	<b>6.0E5</b>	0.017	<b>-5.8</b>
D4A0L4	Y-box-binding protein 3	D4a0l4	5.6E6	4.6E6	8.8E6	<b>6.4E6</b>	0.0E0	1.9E6	0.0E0	3.5E6	0.0E0	<b>1.1E6</b>	0.007	<b>-5.8</b>
G3V6S1	PRKC apoptosis WT1 regulator protein	G3v6s1	2.9E6	3.5E6	1.6E6	<b>2.7E6</b>	9.2E5	0.0E0	1.3E6	0.0E0	0.0E0	<b>4.5E5</b>	0.007	<b>-6.0</b>
G3V6P6	RNA binding motif protein 3, isoform CRA_a	G3v6p6	3.4E7	2.8E7	1.8E7	<b>2.7E7</b>	1.2E7	0.0E0	0.0E0	8.7E6	0.0E0	<b>4.2E6</b>	0.004	<b>-6.4</b>
D3ZFY8	Ubiquitin-conjugating enzyme E2 variant 1-like	D3zfy8	8.3E6	9.6E6	2.3E6	<b>6.7E6</b>	5.8E5	0.0E0	0.0E0	3.7E6	0.0E0	<b>8.6E5</b>	0.022	<b>-7.8</b>
F1LPR6	Immunoglobulin heavy constant mu	F1lpr6	1.5E7	1.6E7	2.5E7	<b>1.9E7</b>	3.0E6	0.0E0	0.0E0	4.9E6	2.7E6	<b>2.1E6</b>	0.001	<b>-8.9</b>
Q6AZ26	C-terminal binding protein 1	Q6az26	3.6E6	2.4E6	2.0E6	<b>2.7E6</b>	1.4E6	0.0E0	0.0E0	0.0E0	0.0E0	<b>2.7E5</b>	0.003	<b>-9.9</b>
P62275	40S ribosomal protein S29	Rs29	8.6E6	7.7E6	3.5E6	<b>6.6E6</b>	3.2E6	0.0E0	0.0E0	0.0E0	0.0E0	<b>6.3E5</b>	0.006	<b>-10.4</b>
B5DF03	Peptidylprolyl isomerase (Fragment)	B5df03	3.2E6	4.1E6	1.9E6	<b>3.1E6</b>	0.0E0	0.0E0	1.5E6	0.0E0	0.0E0	<b>2.9E5</b>	0.004	<b>-10.5</b>
B0BNA7	Eukaryotic translation initiation factor 3 subunit I	Eif3i	3.0E6	3.9E6	2.0E6	<b>3.0E6</b>	0.0E0	1.4E6	0.0E0	0.0E0	0.0E0	<b>2.8E5</b>	0.003	<b>-10.7</b>
Q5XIC1	Mannose-1-phosphate guanyltransferase alpha	Gmppa	2.3E6	3.5E6	0.0E0	<b>1.9E6</b>	0.0E0	0.0E0	0.0E0	0.0E0	0.0E0	<b>0.0E0</b>	0.043	<b>-1.9E6</b>
B2RZA9	Ube2l3 protein	B2rza9	5.0E6	4.3E6	0.0E0	<b>3.1E6</b>	0.0E0	0.0E0	0.0E0	0.0E0	0.0E0	<b>0.0E0</b>	0.035	<b>-3.1E6</b>
A0A140TAA4	Programmed cell death 6-interacting protein	A0a140taa4	3.9E6	4.3E6	2.1E6	<b>3.4E6</b>	0.0E0	0.0E0	0.0E0	0.0E0	0.0E0	<b>0.0E0</b>	0.001	<b>-3.4E6</b>
M0RD20	Calpain small subunit 1	M0rd20	5.6E6	5.0E6	0.0E0	<b>3.5E6</b>	0.0E0	0.0E0	0.0E0	0.0E0	0.0E0	<b>0.0E0</b>	0.034	<b>-3.5E6</b>
Q63002	Mannose 6-phosphate/insulin-like growth factor II receptor	Q63002	5.4E6	5.2E6	0.0E0	<b>3.5E6</b>	0.0E0	0.0E0	0.0E0	0.0E0	0.0E0	<b>0.0E0</b>	0.034	<b>-3.5E6</b>
Q4KM65	Cleavage and polyadenylation specificity factor subunit 5	Cpsf5	6.7E6	5.7E6	0.0E0	<b>4.1E6</b>	0.0E0	0.0E0	0.0E0	0.0E0	0.0E0	<b>0.0E0</b>	0.035	<b>-4.1E6</b>