Table S1. Differentially expressed proteins.

Accession	Protein description	RatioD/F	P value	Gene name	D.D2	D.D4	D.D3	D.D1	F.F2	F.F4	F.F3	F.F1
Q16706	Alpha-mannosidase 2 OS	1.21	0.027	MAN2A1	1.15	1.04	0.91	1.1	0.77	0.91	0.91	0.886
P80723	Brain acid soluble protein	0.758	0.007	BASP1	0.87	0.96	1.06	0.83	1.3	1.23	1.08	1.297
Q13449	Limbic system-associated	0.826	0.042	LSAMP	0.82	0.95	1.01	1.14	1.31	1.15	1.14	1.143
P01023	Alpha-2-macroglobulin O	0.712	0.004	A2M	0.87	0.95	1	1.18	1.31	1.53	1.32	1.466
P20815	Cytochrome P450 3A5 O	0.66	0.004	CYP3A5	0.71		0.73	0.64	1.1		1.12	0.93
P02748	Complement component	0.828	0.01	C9	0.89	1.04	0.87	1.05	1.21	1.16	1.14	1.137
Q9NZP8	Complement C1r subcom	0.811	0.004	C1RL	0.86	0.93	0.99	1.02	1.14	1.13	1.27	1.137
P02765	Alpha-2-HS-glycoprotein	0.828	0.008	AHSG	0.96	0.9	0.85	0.97	1.01	1.08	1.15	1.192
Q9HAT2	Sialate O-acetylesterase O	0.821	0.018	SIAE	0.84	0.86	0.79		1.01	0.93	1.09	
O14498	Immunoglobulin superfar	1.219	0.003	ISLR	1.06	1.12	1.08	1.12	0.9	0.84	0.85	0.998
Q9BS26	Endoplasmic reticulum re	1.432	0.003	ERP44	1.18	1.11	1.25	1.07	0.73	0.77	0.75	0.976
O00602	Ficolin-1 OS=Homo sapi	0.714	0.02	FCN1	0.72	0.96	1.13	0.79	1.42	1.24	1.26	1.132
P11021	Endoplasmic reticulum cl	0.833	0.006	HSPA5	0.95	0.98	0.91	1.04	1.25	1.15	1.18	1.072
P54108	Cysteine-rich secretory pr	0.786	0.005	CRISP3	0.83	0.93	1.01	1	1.28	1.13	1.24	1.132
Q9P121	Neurotrimin OS=Homo s	0.796	0.041	NTM			0.95	0.97			1.26	1.145
P00734	Prothrombin OS=Homo s	1.234	4E-04	F2	1.03	1.06	1.06	1.08	0.92	0.83	0.87	0.805
Q92484	Acid sphingomyelinase-li	0.781	0.022	SMPDL3A	0.91	0.87		0.85	1	1.12		1.245
P14209	CD99 antigen OS=Homo	0.798	0.003	CD99	0.87		0.92	0.84	1.1		1.14	1.053
P01871	Immunoglobulin heavy c	0.819	0.039	IGHM	0.72	0.65	0.72	0.69	0.95	0.83	0.92	0.695
P01709	Immunoglobulin lambda	0.753	0.033	IGLV2-8		0.73	0.66	0.78		0.98	1.07	0.833
P00558	Phosphoglycerate kinase	0.721	0.016	PGK1	0.72	0.8	0.84	1.04	1.4	1.08	1.1	1.125
Q9NZK5	Adenosine deaminase 2 O	0.805	0.002	ADA2	0.94	0.98	0.89	0.88	1.27	1.13	1.11	1.079
Q96RT7	Gamma-tubulin complex	1.549	0.008	TUBGCP6		1.17		1.1		0.75		0.718
Q86VW0	SEC14 domain and spect	0.53	2E-04	SESTD1	0.57	0.8	0.66	0.64	1.29	1.25	1.33	1.159
P08670	Vimentin OS=Homo sapi	0.362	0.02	VIM	0.64			0.82	2.19			1.845
P02776	Platelet factor 4 OS=Hon	2.877	0.018	PF4	2.13		1.01	1.15	0.58		0.38	0.53
P14151	L-selectin OS=Homo sap	0.764	0.007	SELL	0.86	0.96	0.98	1.07	1.45	1.24	1.25	1.143
Q15149	Plectin OS=Homo sapien	0.545	0.036	PLEC	0.57			0.7	1.23			1.101
O60832	H/ACA ribonucleoprotein	1.252	0.037	DKC1		0.89		0.95		0.76		0.715
P07602	Prosaposin OS=Homo sa	0.831	0.019	PSAP	0.79	0.86	1.02	0.98	1.08	1.09	1.15	1.077
O00461	Golgi integral membrane	0.77	0.028	GOLIM4	0.99		0.9		1.23		1.22	
P15941	Mucin-1 OS=Homo sapie	1.36	0.01	MUC1		1.15		1.21		0.85		0.878
Q86YW5	Trem-like transcript 1 pro	0.826	0.007	TREML1	0.86	0.8	0.73	0.87	0.98	0.97	1.06	0.937
P02679	Fibrinogen gamma chain	4.174	0.013	FGG	2.65	1.06	0.88	1.56	0.42	0.36	0.35	0.341
P09486	SPARC OS=Homo sapie	0.821	0.003	SPARC	0.98	0.9	0.9	0.93	1.08	1.11	1.25	1.069
P10619	Lysosomal protective pro	0.64	0.032	CTSA		0.83		0.93		1.3		1.458
P02675	Fibrinogen beta chain OS	4.694	0.002	FGB	2.48	1.09	0.88	1.6	0.41	0.23	0.41	0.249
P13489	Ribonuclease inhibitor O	0.806	0.037	RNH1		0.92	1.01	1.05		1.18	1.37	1.14
A0A075B	Immunoglobulin lambda	0.613	0.004	IGLV3-9	0.65	0.53	0.77	0.61	1.07	1.07	1.2	0.833
P14618	Pyruvate kinase PKM OS	0.829	0.002	PKM	0.96	0.96	0.9	1	1.16	1.07	1.23	1.149
P02647	Apolipoprotein A-I OS=H	0.797	0.024	APOA1	0.84	0.99	1.15	0.97	1.35	1.23	1.27	1.111
P29122	Proprotein convertase sul	0.829	0.02	PCSK6	0.87	0.89	0.94		1.06	1.17	1.01	
Q8IUL8	Cartilage intermediate lay	0.652	4E-04	CILP2	1.03	0.93	0.96	0.85	1.57	1.33	1.54	1.345
Q9H299	SH3 domain-binding glut	0.658	0.049	SH3BGRL3	0.68	0.58	0.69	0.86	1.25	0.97	1.33	0.714
P04085	Platelet-derived growth fa	0.571	0.002	PDGFA	0.68	0.88	0.78	0.93	1.53	1.37	1.67	1.156
P18428	Lipopolysaccharide-bindi	1.703	0.038	LBP		1.18	1.36	0.95		0.76	0.5	0.786
O76074	cGMP-specific 3',5'-cycli	1.528	0.035	PDE5A	1.1		1.19		0.7		0.8	
Q8N3X6	Ligand-dependent nuclea	0.772	0.017	LCORL		0.96	0.84	1.03		1.18	1.3	1.184
P22304	Iduronate 2-sulfatase OS=	0.823	0.047	IDS	0.84	0.9	1.05		1.18	1.11	1.1	
P0DOY3	Immunoglobulin lambda	0.774	0.013	IGLC3	0.75	0.76	0.91	0.85	1.13	1.06	1.15	0.895
P04114	Apolipoprotein B-100 OS	1.744	0.01	APOB	1.35	1.11	0.83	0.95	0.47	0.8	0.59	0.566

Q9ULV0	Unconventional myosin-V	0.727	0.043	MYO5B		0.8	0.77			1.01	1.15	
P02775	Platelet basic protein OS=	0.735	0.007	PPBP	0.81	0.9	0.84	1.03	1.18	1.17	1.43	1.088
P02671	Fibrinogen alpha chain O	2.197	0.003	FGA	1.51	1.01	0.87	1.4	0.7	0.53	0.48	0.474
P09972	Fructose-bisphosphate al	0.776	0.031	ALDOC	0.95		0.87		1.19		1.16	
P08709	Coagulation factor VII O	0.757	0.018	F7	0.75	0.87		0.92	1.04	1.14		1.179
P0C0L4	Complement C4-A OS=H	1.342	0.004	C4A	1.24	1.11	1.42	1.12	0.96	0.93	0.84	0.918
P0C0L5	Complement C4-B OS=H	1.298	0.001	C4B	1.21	1.06	1	1.14	0.86	0.86	0.88	0.799
Q13201	Multimerin-1 OS=Homo	0.796	0.001	MMRN1	0.87	0.92	0.92	1.01	1.22	1.19	1.17	1.093
P05154	Plasma serine protease in	0.81	0.049	SERPINA5	0.88	0.89	0.8	1.09	1.14	1.15	1.08	1.15
P05067	Amyloid-beta precursor p	0.707	2E-05	APP	0.85	0.88	0.87	0.92	1.21	1.19	1.27	1.303
P00738	Haptoglobin OS=Homo s	1.605	0.041	HP	1.42	1.29	1.7	0.84	0.87	0.83	0.95	0.628
P00742	Coagulation factor X OS=	0.812	0.024	F10	0.93	0.91	0.92	0.95	1.02	1.09	1.22	1.252
P24298	Alanine aminotransferase	0.808	0.018	GPT	0.92	0.92			1.17	1.11		
Q14956	Transmembrane glycoprc	1.316	0.027	GPNMB	1.32	1.18	1.02	1.32	0.81	0.93	0.83	1.114
P62805	Histone H4 OS=Homo sa	0.642	0.018	HIST1H4A	0.93	0.68	0.73	0.74	1.49	1.18	0.86	1.254
Q9Y5C1	Angiopoietin-related prot	0.817	0.019	ANGPTL3	0.74	0.88	0.9	0.92	1.15	1.04	1.08	0.944
P07738	Bisphosphoglycerate mut	0.677	0.042	BPGM	0.6	0.93	0.75	1.13	1.22	1.17	1.57	1.067