Electronic Supplemental Material Table 1

The Table lists proteins that were identified in this study. Accession number and protein names are shown. AS 116/114, 116/115, 117/114, 117/115 are the internal iTRAQ technical replicates, while AS 115/114 and 117/116 are the internal swap technical controls. Pink-shaded cells show proteins downregulated in non-healing (NH) compared to rapidly healing (RH) wounds. Green-shaded cells show proteins upregulated in NH vs RH wounds. SerpinB3 is underlined.

Accession	Description	A5: 115/114	A5: 117/116	A5: 116/114	A5: 116/115	A5: 117/114	A5: 117/115	Average NH / RH ratio
P20930	Filaggrin OS=Homo sapiens GN=FLG PE=1 SV=3 - [FILA_HUMAN]	1.28	1.11	0.27	0.12	0.33	0.19	0.23
P21810	Biglycan OS=Homo sapiens GN=BGN PE=1 SV=2 - [PGS1_HUMAN]	1.01	1.17	0.33	0.32	0.37	0.37	0.35
P51884	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2 - [LUM_HUMAN]	1.04	1.08	0.36	0.35	0.39	0.39	0.37
P08123	Collagen alpha-2(I) chain OS=Homo sapiens GN=COL1A2 PE=1 SV=7 - [CO1A2_HUMAN]	0.99	1.36	0.41	0.45	0.57	0.64	0.52
P31944	Caspase-14 OS=Homo sapiens GN=CASP14 PE=1 SV=2 - [CASPE_HUMAN]	1.04	1.05	0.41	0.41	0.45	0.42	0.42
P26373	60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4 - [RL13_HUMAN]	1.09	0.96	0.43	0.39	0.42	0.38	0.40
P47929	Galectin-7 OS=Homo sapiens GN=LGALS7 PE=1 SV=2 - [LEG7_HUMAN]	0.95	1.06	0.44	0.53	0.43	0.52	0.48
P02452	Collagen alpha-1(I) chain OS=Homo sapiens GN=COL1A1 PE=1 SV=5 - [CO1A1_HUMAN]	1.00	1.14	0.44	0.45	0.63	0.55	0.52
Q15063-3	Isoform 3 of Periostin OS=Homo sapiens GN=POSTN - [POSTN_HUMAN]	1.01	0.99	0.45	0.39	0.51	0.46	0.45
Q6UWP8	Suprabasin OS=Homo sapiens GN=SBSN PE=2 SV=1 - [SBSN_HUMAN]	1.07	1.04	0.47	0.44	0.48	0.44	0.46
P24534	Elongation factor 1-beta OS=Homo sapiens GN=EEF1B2 PE=1 SV=3 - [EF1B_HUMAN]	1.03	1.14	0.48	0.46	0.57	0.54	0.51
P31151	Protein S100-A7 OS=Homo sapiens GN=S100A7 PE=1 SV=4 - [S10A7_HUMAN]	1.08	0.98	0.48	0.44	0.46	0.42	0.45
Q5VTM1	Protein FAM25 OS=Homo sapiens GN=FAM25A PE=2 SV=2 - [FAM25_HUMAN]	1.03	1.06	0.49	0.50	0.50	0.47	0.49
P19957	Elafin OS=Homo sapiens GN=PI3 PE=1 SV=3 - [ELAF_HUMAN]	1.03	1.05	0.49	0.46	0.48	0.47	0.48
P54652	Heat shock-related 70 kDa protein 2 OS=Homo sapiens GN=HSPA2 PE=1 SV=1 - [HSP72_HUMAN]	0.99	1.10	0.49	0.49	0.52	0.52	0.51
Q8NC51-4	Isoform 4 of Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERBP1 - [PAIRB_HUMAN]	0.93	1.10	0.49	0.52	0.46	0.54	0.50
P20810-5	Isoform 5 of Calpastatin OS=Homo sapiens GN=CAST - [ICAL_HUMAN]	0.96	0.94	0.49	0.50	0.46	0.48	0.48
Q08554-2	Isoform 1B of Desmocollin-1 OS=Homo sapiens GN=DSC1 - [DSC1_HUMAN]	0.86	0.86	0.50	0.58	0.42	0.48	0.49
Q6E0U4-5	Isoform 5 of Dermokine OS=Homo sapiens GN=DMKN - [DMKN_HUMAN]	1.03	1.08	0.50	0.47	0.48	0.43	0.47
P05089	Arginase-1 OS=Homo sapiens GN=ARG1 PE=1 SV=2 - [ARGI1_HUMAN]	0.97	1.04	0.51	0.45	0.53	0.52	0.50
O14737	Programmed cell death protein 5 OS=Homo sapiens GN=PDCD5 PE=1 SV=3 - [PDCD5_HUMAN]	1.10	1.01	0.51	0.46	0.50	0.45	0.48
P27816-6	Isoform 6 of Microtubule-associated protein 4 OS=Homo sapiens GN=MAP4 - [MAP4_HUMAN]	0.95	1.05	0.52	0.48	0.52	0.49	0.50
P06748-2	Isoform 2 of Nucleophosmin OS=Homo sapiens GN=NPM1 - [NPM_HUMAN]	0.98	1.02	0.52	0.51	0.51	0.52	0.52
P23588	Eukaryotic translation initiation factor 4B OS=Homo sapiens GN=EIF4B PE=1 SV=2 - [IF4B_HUMAN]	1.01	1.10	0.55	0.66	0.60	0.71	0.63
P05387	60S acidic ribosomal protein P2 OS=Homo sapiens GN=RPLP2 PE=1 SV=1 - [RLA2_HUMAN]	1.03	1.02	0.55	0.55	0.72	0.80	0.65
P29508	**Serpin B3 OS=Homo sapiens GN=SERPINB3 PE=1 SV=2 - [SPB3_HUMAN]**	**1.01**	**1.04**	0.55	0.53	0.58	0.54	**0.55**
P19105	Myosin regulatory light chain 12A OS=Homo sapiens GN=MYL12A PE=1 SV=2 - [ML12A_HUMAN]	0.99	1.15	0.56	0.56	0.56	0.59	0.57
P25311	Zinc-alpha-2-glycoprotein OS=Homo sapiens GN=AZGP1 PE=1 SV=2 - [ZA2G_HUMAN]	1.07	1.00	0.56	0.58	0.60	0.58	0.58
P37802	Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3 - [TAGL2_HUMAN]	0.98	1.02	0.56	0.57	0.57	0.57	0.57
P63167	Dynein light chain 1, cytoplasmic OS=Homo sapiens GN=DYNLL1 PE=1 SV=1 - [DYL1_HUMAN]	1.09	1.12	0.56	0.51	0.61	0.56	0.56
Q9BW30	Tubulin polymerization-promoting protein family member 3 OS=Homo sapiens GN=TPPP3 PE=1 SV=1 - [redacted]	0.89	0.97	0.57	0.58	0.61	0.54	0.58
Q5D862	Filaggrin-2 OS=Homo sapiens GN=FLG2 PE=1 SV=1 - [FILA2_HUMAN]	0.93	0.91	0.57	0.61	0.51	0.54	0.56
P63241	Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A PE=1 SV=2 - [IF5A1_HUMAN]	1.00	1.00	0.57	0.58	0.57	0.55	0.57
P67936-2	Isoform 2 of Tropomyosin alpha-4 chain OS=Homo sapiens GN=TPM4 - [TPM4_HUMAN]	0.97	0.99	0.58	0.57	0.57	0.55	0.57
P06702	Protein S100-A9 OS=Homo sapiens GN=S100A9 PE=1 SV=1 - [S10A9_HUMAN]	1.03	0.94	0.58	0.57	0.59	0.55	0.57
Q14134-2	Isoform Beta of Tripartite motif-containing protein 29 OS=Homo sapiens GN=TRIM29 - [TRI29_HUMAN]	0.94	1.00	0.58	0.61	0.57	0.59	0.59
P62750	60S ribosomal protein L23a OS=Homo sapiens GN=RPL23A PE=1 SV=1 - [RL23A_HUMAN]	1.03	0.99	0.59	0.57	0.56	0.54	0.57
Q9BZJ3	Tryptase delta OS=Homo sapiens GN=TPSD1 PE=1 SV=2 - [TRYD_HUMAN]	0.97	1.07	0.60	0.59	0.62	0.63	0.61
P29966	Myristoylated alanine-rich C-kinase substrate OS=Homo sapiens GN=MARCKS PE=1 SV=4 - [MARCS_HUMAN]	0.98	1.00	0.60	0.64	0.58	0.63	0.61
O00515	Ladinin-1 OS=Homo sapiens GN=LAD1 PE=1 SV=2 - [LAD1_HUMAN]	1.01	1.04	0.61	0.53	0.62	0.54	0.57
P27482	Calmodulin-like protein 3 OS=Homo sapiens GN=CALML3 PE=1 SV=2 - [CALL3_HUMAN]	1.02	1.04	0.62	0.62	0.62	0.62	0.62
Q13765	Nascent polypeptide-associated complex subunit alpha OS=Homo sapiens GN=NACA PE=1 SV=1 - [NACA_HUMAN]	1.05	0.99	0.62	0.53	0.59	0.62	0.59
Q9H299	SH3 domain-binding glutamic acid-rich-like protein 3 OS=Homo sapiens GN=SH3BGRL3 PE=1 SV=1 - [SH3L3_HUMAN]	1.00	1.02	0.63	0.60	0.60	0.60	0.61
P60660	Myosin light polypeptide 6 OS=Homo sapiens GN=MYL6 PE=1 SV=2 - [MYL6_HUMAN]	1.00	0.99	0.64	0.63	0.61	0.61	0.62
P16989-2	Isoform 2 of DNA-binding protein A OS=Homo sapiens GN=CSDA - [DBPA_HUMAN]	1.01	1.05	0.64	0.62	0.64	0.63	0.63
P62158	Calmodulin OS=Homo sapiens GN=CALM1 PE=1 SV=2 - [CALM_HUMAN]	0.98	0.94	0.65	0.60	0.58	0.55	0.59
P10599	Thioredoxin OS=Homo sapiens GN=TXN PE=1 SV=3 - [THIO_HUMAN]	1.08	0.97	0.65	0.64	0.65	0.62	0.64

P11940-2	Isoform 2 of Polyadenylate-binding protein 1 OS=Homo sapiens GN=PABPC1 - [PABP1_HUMAN]	0.88	1.07	**0.66**	0.74	0.67	0.77	0.71
Q04760-2	Isoform 2 of Lactoylglutathione lyase OS=Homo sapiens GN=GLO1 - [LGUL_HUMAN]	0.99	1.05	0.67	0.71	0.72	0.69	0.70
P07476	Involucrin OS=Homo sapiens GN=IVL PE=1 SV=2 - [INVO_HUMAN]	1.05	1.00	0.67	**0.64**	**0.64**	**0.58**	0.63
P06753-2	Isoform 2 of Tropomyosin alpha-3 chain OS=Homo sapiens GN=TPM3 - [TPM3_HUMAN]	1.00	1.01	0.67	0.68	**0.64**	0.67	0.67
Q8IZP2	Putative protein FAM10A4 OS=Homo sapiens GN=ST13P4 PE=5 SV=1 - [ST134_HUMAN]	0.94	1.08	0.68	0.72	0.77	0.81	0.74
P09497-2	Isoform Non-brain of Clathrin light chain B OS=Homo sapiens GN=CLTB - [CLCB_HUMAN]	1.00	0.99	0.68	0.68	**0.65**	**0.65**	0.67
P20962	Parathymosin OS=Homo sapiens GN=PTMS PE=1 SV=2 - [PTMS_HUMAN]	0.83	1.00	0.68	0.81	**0.55**	**0.60**	0.66
P31947	14-3-3 protein sigma OS=Homo sapiens GN=SFN PE=1 SV=1 - [1433S_HUMAN]	1.02	1.02	0.70	0.68	0.71	0.68	0.69
P05109	Protein S100-A8 OS=Homo sapiens GN=S100A8 PE=1 SV=1 - [S10A8_HUMAN]	1.11	1.00	0.70	**0.60**	**0.66**	**0.59**	0.64
Q9NZT1	Calmodulin-like protein 5 OS=Homo sapiens GN=CALML5 PE=1 SV=2 - [CALL5_HUMAN]	1.07	1.00	0.70	0.71	0.69	0.70	0.70
P09382	Galectin-1 OS=Homo sapiens GN=LGALS1 PE=1 SV=2 - [LEG1_HUMAN]	0.91	1.05	0.70	0.67	0.68	**0.66**	0.68
O75347	Tubulin-specific chaperone A OS=Homo sapiens GN=TBCA PE=1 SV=3 - [TBCA_HUMAN]	0.94	1.03	0.70	0.74	0.69	0.73	0.72
P16403	Histone H1.2 OS=Homo sapiens GN=HIST1H1C PE=1 SV=2 - [H12_HUMAN]	1.00	0.96	0.71	0.73	**0.66**	0.68	0.69
P06454-2	Isoform 2 of Prothymosin alpha OS=Homo sapiens GN=PTMA - [PTMA_HUMAN]	1.05	0.97	0.72	0.67	0.69	0.69	0.69
P51858	Hepatoma-derived growth factor OS=Homo sapiens GN=HDGF PE=1 SV=1 - [HDGF_HUMAN]	1.04	0.92	0.72	0.75	**0.63**	**0.66**	0.69
P28070	Proteasome subunit beta type-4 OS=Homo sapiens GN=PSMB4 PE=1 SV=4 - [PSB4_HUMAN]	0.67	1.07	0.72	0.90	0.90	1.06	0.90
P19971	Thymidine phosphorylase OS=Homo sapiens GN=TYMP PE=1 SV=2 - [TYPH_HUMAN]	0.89	1.18	0.72	0.80	0.82	0.92	0.82
Q09666	Neuroblast differentiation-associated protein AHNAK OS=Homo sapiens GN=AHNAK PE=1 SV=2 - [AHNK_HUMAN]	0.94	0.99	0.72	0.72	0.72	0.72	0.72
P07910-3	Isoform 3 of Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC - [HNRPC_HUMAN]	0.99	1.10	0.72	0.76	0.72	0.76	0.74
P19338	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3 - [NUCL_HUMAN]	0.98	1.02	0.72	0.72	0.71	0.74	0.72
P61604	10 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPE1 PE=1 SV=2 - [CH10_HUMAN]	1.04	1.06	0.73	0.68	0.71	**0.63**	0.69
P39019	40S ribosomal protein S19 OS=Homo sapiens GN=RPS19 PE=1 SV=2 - [RS19_HUMAN]	0.93	1.02	0.73	0.77	0.67	0.72	0.72
P67936	Tropomyosin alpha-4 chain OS=Homo sapiens GN=TPM4 PE=1 SV=3 - [TPM4_HUMAN]	1.04	0.98	0.74	0.71	0.71	0.69	0.72
P22626-2	Isoform A2 of Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 - [ROA2_HUMAN]	1.02	1.01	0.74	0.76	0.71	0.74	0.74
P80723	Brain acid soluble protein 1 OS=Homo sapiens GN=BASP1 PE=1 SV=2 - [BASP1_HUMAN]	0.89	1.12	0.75	0.95	0.86	0.93	0.87
P54727	UV excision repair protein RAD23 homolog B OS=Homo sapiens GN=RAD23B PE=1 SV=1 - [RD23B_HUMAN]	0.97	1.05	0.75	0.78	0.76	0.73	0.76
P11021	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2 - [GRP78_HUMAN]	0.94	0.97	0.76	0.82	0.81	0.83	0.80
P31949	Protein S100-A11 OS=Homo sapiens GN=S100A11 PE=1 SV=2 - [S10AB_HUMAN]	1.01	1.07	0.76	0.69	0.74	0.70	0.72
Q92820	Gamma-glutamyl hydrolase OS=Homo sapiens GN=GGH PE=1 SV=2 - [GGH_HUMAN]	0.94	0.92	0.76	0.80	0.67	0.71	0.74
P29034	Protein S100-A2 OS=Homo sapiens GN=S100A2 PE=1 SV=3 - [S10A2_HUMAN]	1.08	0.93	0.76	0.70	0.67	**0.62**	0.69
P09622	Dihydrolipoyl dehydrogenase, mitochondrial OS=Homo sapiens GN=DLD PE=1 SV=2 - [DLDH_HUMAN]	0.92	0.98	0.78	0.83	0.67	0.72	0.75
P27797	Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1 - [CALR_HUMAN]	0.97	1.00	0.79	0.80	0.78	0.75	0.78
P60953	Cell division control protein 42 homolog OS=Homo sapiens GN=CDC42 PE=1 SV=2 - [CDC42_HUMAN]	0.84	0.98	0.79	0.93	0.74	0.88	0.83
P28066	Proteasome subunit alpha type-5 OS=Homo sapiens GN=PSMA5 PE=1 SV=3 - [PSA5_HUMAN]	0.94	0.99	0.79	0.77	0.80	0.79	0.78
P35241	Radixin OS=Homo sapiens GN=RDX PE=1 SV=1 - [RADI_HUMAN]	0.83	1.25	0.79	0.93	0.94	1.13	0.95
P40121	Macrophage-capping protein OS=Homo sapiens GN=CAPG PE=1 SV=2 - [CAPG_HUMAN]	0.95	1.01	0.79	0.82	0.77	0.80	0.79
P51991-2	Isoform 2 of Heterogeneous nuclear ribonucleoprotein A3 OS=Homo sapiens GN=HNRNPA3 - [ROA3_HUMAN]	1.08	1.13	0.79	0.73	0.86	0.79	0.79
P62937	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2 - [PPIA_HUMAN]	1.05	1.01	0.79	0.72	0.79	0.76	0.77
P09493-4	Isoform 4 of Tropomyosin alpha-1 chain OS=Homo sapiens GN=TPM1 - [TPM1_HUMAN]	0.86	0.90	0.79	0.91	0.69	0.79	0.80
P30101	Protein disulfide-isomerase A3 OS=Homo sapiens GN=PDIA3 PE=1 SV=4 - [PDIA3_HUMAN]	1.03	1.05	0.80	0.76	0.99	0.94	0.87
P17900	Ganglioside GM2 activator OS=Homo sapiens GN=GM2A PE=1 SV=4 - [SAP3_HUMAN]	0.93	1.03	0.80	0.85	0.90	0.96	0.88
P07355	Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2 - [ANXA2_HUMAN]	1.02	1.00	0.80	0.77	0.75	0.72	0.76
P63104	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1 - [1433Z_HUMAN]	1.01	1.01	0.80	0.82	0.93	0.86	0.85
Q01105-2	Isoform 2 of Protein SET OS=Homo sapiens GN=SET - [SET_HUMAN]	1.11	1.01	0.80	0.72	0.91	0.80	0.81
A8MW06	Thymosin beta-4-like protein 3 OS=Homo sapiens GN=TMSL3 PE=2 SV=1 - [TMSL3_HUMAN]	1.02	1.01	0.81	0.72	0.77	0.70	0.75
P22352	Glutathione peroxidase 3 OS=Homo sapiens GN=GPX3 PE=1 SV=2 - [GPX3_HUMAN]	0.99	0.97	0.81	0.79	0.76	0.76	0.78
P22307-2	Isoform SCP2 of Non-specific lipid-transfer protein OS=Homo sapiens GN=SCP2 - [NLTP_HUMAN]	0.99	1.08	0.81	0.81	0.84	0.85	0.83
O75368	SH3 domain-binding glutamic acid-rich-like protein OS=Homo sapiens GN=SH3BGRL PE=1 SV=1 - [SH3L1_HUMAN]	1.02	1.09	0.81	0.79	0.83	0.77	0.80

Q9NQ38	Serine protease inhibitor Kazal-type 5 OS=Homo sapiens GN=SPINK5 PE=1 SV=2 - [ISK5_HUMAN]	0.99	1.02	0.81	0.76	0.79	0.76	0.78
P20774	Mimecan OS=Homo sapiens GN=OGN PE=1 SV=1 - [MIME_HUMAN]	1.05	1.35	0.81	0.76	**0.36**	**0.35**	0.57
P46940	Ras GTPase-activating-like protein IQGAP1 OS=Homo sapiens GN=IQGAP1 PE=1 SV=1 - [IQGA1_HUMAN]	1.02	0.97	0.82	0.79	0.77	0.75	0.78
P22528	Cornifin-B OS=Homo sapiens GN=SPRR1B PE=1 SV=2 - [SPR1B_HUMAN]	0.90	0.88	0.82	0.91	0.70	0.77	0.80
P04792	Heat shock protein beta-1 OS=Homo sapiens GN=HSPB1 PE=1 SV=2 - [HSPB1_HUMAN]	1.04	1.00	0.83	0.79	0.85	0.73	0.80
P14314	Glucosidase 2 subunit beta OS=Homo sapiens GN=PRKCSH PE=1 SV=2 - [GLU2B_HUMAN]	1.00	1.04	0.83	0.82	0.80	0.80	0.81
P35908	Keratin, type II cytoskeletal 2 epidermal OS=Homo sapiens GN=KRT2 PE=1 SV=2 - [K22E_HUMAN]	1.03	0.99	0.83	0.80	0.79	0.78	0.80
P12111-2	Isoform 2 of Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 - [CO6A3_HUMAN]	1.03	1.01	0.83	0.81	0.95	0.93	0.88
Q9H0W9-3	Isoform 3 of Ester hydrolase C11orf54 OS=Homo sapiens GN=C11orf54 - [CK054_HUMAN]	0.74	0.97	0.83	1.12	0.77	1.05	0.94
P13645	Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6 - [K1C10_HUMAN]	1.11	1.02	0.84	0.70	0.81	0.72	0.77
P01714	Ig lambda chain V-III region SH OS=Homo sapiens PE=1 SV=1 - [LV301_HUMAN]	1.00	0.87	0.84	0.83	0.71	0.70	0.77
P11279	Lysosome-associated membrane glycoprotein 1 OS=Homo sapiens GN=LAMP1 PE=1 SV=3 - [LAMP1_HUMAN]	1.02	1.09	0.84	0.82	0.88	0.86	0.85
O75369-7	Isoform 7 of Filamin-B OS=Homo sapiens GN=FLNB - [FLNB_HUMAN]	0.96	1.09	0.84	0.88	0.86	0.90	0.87
Q86UP3-3	Isoform 3 of Zinc finger homeobox protein 4 OS=Homo sapiens GN=ZFHX4 - [ZFHX4_HUMAN]	0.84	0.52	0.85	1.00	**0.42**	**0.50**	0.69
P22392-2	Isoform 3 of Nucleoside diphosphate kinase B OS=Homo sapiens GN=NME2 - [NDKB_HUMAN]	1.00	1.01	0.85	0.81	0.83	0.81	0.82
P24821-4	Isoform 4 of Tenascin OS=Homo sapiens GN=TNC - [TENA_HUMAN]	1.00	1.01	0.86	0.82	0.84	0.84	0.84
Q99436	Proteasome subunit beta type-7 OS=Homo sapiens GN=PSMB7 PE=1 SV=1 - [PSB7_HUMAN]	1.11	1.06	0.86	0.77	0.88	0.79	0.82
P38159-2	Isoform 2 of RNA-binding motif protein, X chromosome OS=Homo sapiens GN=RBMX - [RBMX_HUMAN]	0.90	0.91	0.86	0.95	0.79	0.93	0.88
P37837	Transaldolase OS=Homo sapiens GN=TALDO1 PE=1 SV=2 - [TALDO_HUMAN]	0.87	1.05	0.86	1.24	1.00	1.25	1.09
P20290-2	Isoform 2 of Transcription factor BTF3 OS=Homo sapiens GN=BTF3 - [BTF3_HUMAN]	1.14	0.90	0.86	0.78	0.75	0.68	0.77
O14818	Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1 - [PSA7_HUMAN]	0.94	1.08	0.87	0.92	0.88	0.94	0.90
P48637	Glutathione synthetase OS=Homo sapiens GN=GSS PE=1 SV=1 - [GSHB_HUMAN]	0.99	1.13	0.87	0.81	0.73	0.74	0.79
Q01469	Fatty acid-binding protein, epidermal OS=Homo sapiens GN=FABP5 PE=1 SV=3 - [FABP5_HUMAN]	1.02	1.00	0.87	0.83	0.84	0.81	0.84
Q06323	Proteasome activator complex subunit 1 OS=Homo sapiens GN=PSME1 PE=1 SV=1 - [PSME1_HUMAN]	0.88	0.77	0.88	0.98	**0.65**	0.73	0.81
P31946-2	Isoform Short of 14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB - [1433B_HUMAN]	1.00	0.97	0.89	0.87	0.80	0.79	0.84
P23284	Peptidyl-prolyl cis-trans isomerase B OS=Homo sapiens GN=PPIB PE=1 SV=2 - [PPIB_HUMAN]	0.96	1.04	0.89	0.93	0.89	0.95	0.92
P61916	Epididymal secretory protein E1 OS=Homo sapiens GN=NPC2 PE=1 SV=1 - [NPC2_HUMAN]	0.90	1.06	0.89	0.76	0.84	0.75	0.81
P07237	Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3 - [PDIA1_HUMAN]	1.09	0.91	0.89	0.87	0.82	0.79	0.85
P13473	Lysosome-associated membrane glycoprotein 2 OS=Homo sapiens GN=LAMP2 PE=1 SV=2 - [LAMP2_HUMAN]	1.01	1.03	0.89	0.96	0.83	0.82	0.88
P09211	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2 - [GSTP1_HUMAN]	0.97	1.00	0.90	0.83	0.80	0.81	0.83
Q9ULZ3-2	Isoform 2 of Apoptosis-associated speck-like protein containing a CARD OS=Homo sapiens GN=PYCARD -	0.93	1.06	0.90	1.02	0.90	1.01	0.96
P13987	CD59 glycoprotein OS=Homo sapiens GN=CD59 PE=1 SV=1 - [CD59_HUMAN]	1.01	1.00	0.90	0.88	0.86	0.85	0.87
P07108	Acyl-CoA-binding protein OS=Homo sapiens GN=DBI PE=1 SV=2 - [ACBP_HUMAN]	1.12	1.01	0.90	0.81	0.90	0.82	0.86
P09651-3	Isoform 2 of Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 - [ROA1_HUMAN]	1.12	1.00	0.90	0.79	0.84	0.77	0.83
P04080	Cystatin-B OS=Homo sapiens GN=CSTB PE=1 SV=2 - [CYTB_HUMAN]	1.07	0.98	0.91	0.82	0.79	0.75	0.82
P58107	Epiplakin OS=Homo sapiens GN=EPPK1 PE=1 SV=2 - [EPIPL_HUMAN]	0.96	1.04	0.91	0.92	1.16	1.31	1.07
P11142	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1 - [HSP7C_HUMAN]	1.04	1.09	0.91	0.88	0.91	0.89	0.90
O95274	Ly6/PLAUR domain-containing protein 3 OS=Homo sapiens GN=LYPD3 PE=1 SV=2 - [LYPD3_HUMAN]	1.01	1.00	0.91	0.90	0.88	0.87	0.89
P99999	Cytochrome c OS=Homo sapiens GN=CYCS PE=1 SV=2 - [CYC_HUMAN]	1.00	1.00	0.92	0.96	0.89	0.89	0.92
P51888	Prolargin OS=Homo sapiens GN=PRELP PE=1 SV=1 - [PRELP_HUMAN]	1.08	1.13	0.94	0.86	0.95	0.87	0.91
P14174	Macrophage migration inhibitory factor OS=Homo sapiens GN=MIF PE=1 SV=4 - [MIF_HUMAN]	1.09	1.05	0.95	0.91	0.96	0.90	0.93
P04075	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2 - [ALDOA_HUMAN]	0.98	1.00	0.95	1.00	1.01	1.01	0.99
Q99497	Protein DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2 - [PARK7_HUMAN]	0.95	1.04	0.95	0.99	1.00	1.00	0.99
P06576	ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3 - [ATPB_HUMAN]	0.94	0.97	0.96	1.01	0.90	0.95	0.96
O43399-2	Isoform 2 of Tumor protein D54 OS=Homo sapiens GN=TPD52L2 - [TPD54_HUMAN]	1.06	1.01	0.97	0.91	0.94	0.89	0.92
P15311	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4 - [EZRI_HUMAN]	1.01	0.99	0.97	0.96	0.89	0.86	0.92
P18206-2	Isoform 1 of Vinculin OS=Homo sapiens GN=VCL - [VINC_HUMAN]	0.97	1.03	0.98	1.01	1.00	0.98	0.99
P08758	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2 - [ANXA5_HUMAN]	0.93	0.98	0.99	1.04	0.95	0.99	0.99

P02766	Transthyretin OS=Homo sapiens GN=TTR PE=1 SV=1 - [TTHY_HUMAN]	1.00	1.08	0.99	0.98	0.99	1.01	0.99
P01040	Cystatin-A OS=Homo sapiens GN=CSTA PE=1 SV=1 - [CYTA_HUMAN]	1.08	0.98	1.01	0.94	0.97	0.89	0.95
P0CG47	Polyubiquitin-B OS=Homo sapiens GN=UBB PE=1 SV=1 - [UBB_HUMAN]	1.07	0.97	1.01	0.93	0.95	0.90	0.95
Q13561	Dynactin subunit 2 OS=Homo sapiens GN=DCTN2 PE=1 SV=4 - [DCTN2_HUMAN]	0.92	1.07	1.03	0.86	1.07	1.15	1.03
P68104	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1 - [EF1A1_HUMAN]	1.01	1.00	1.03	0.99	1.00	0.96	1.00
P49720	Proteasome subunit beta type-3 OS=Homo sapiens GN=PSMB3 PE=1 SV=2 - [PSB3_HUMAN]	1.00	1.19	1.05	1.04	1.20	1.19	1.12
P06396-2	Isoform 2 of Gelsolin OS=Homo sapiens GN=GSN - [GELS_HUMAN]	0.94	0.97	1.06	1.08	1.02	1.04	1.05
P25788-2	Isoform 2 of Proteasome subunit alpha type-3 OS=Homo sapiens GN=PSMA3 - [PSA3_HUMAN]	0.96	1.01	1.07	1.10	1.10	1.06	1.08
P60709	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]	1.06	0.97	1.07	1.00	1.08	1.00	1.04
P26038	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3 - [MOES_HUMAN]	1.02	0.96	1.07	1.06	1.01	0.97	1.03
P15924-2	Isoform DPII of Desmoplakin OS=Homo sapiens GN=DSP - [DESP_HUMAN]	1.00	1.02	1.07	1.05	1.07	1.02	1.05
P02545-2	Isoform C of Prelamin-A/C OS=Homo sapiens GN=LMNA - [LMNA_HUMAN]	1.04	0.97	1.07	1.04	1.03	1.00	1.04
Q13011	Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial OS=Homo sapiens GN=ECH1 PE=1 SV=2 - [ECH1_HUMAN]	1.01	0.95	1.07	1.05	0.98	0.97	1.02
P62258	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1 - [1433E_HUMAN]	0.98	1.00	1.08	1.09	1.04	1.06	1.07
P13639	Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4 - [EF2_HUMAN]	0.94	1.08	1.08	1.06	1.33	**1.64**	1.28
P01625	Ig kappa chain V-IV region Len OS=Homo sapiens PE=1 SV=2 - [KV402_HUMAN]	1.09	1.10	1.09	1.01	1.16	1.08	1.09
P04083	Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2 - [ANXA1_HUMAN]	1.02	0.99	1.10	1.10	0.99	1.06	1.06
P35527	Keratin, type I cytoskeletal 9 OS=Homo sapiens GN=KRT9 PE=1 SV=3 - [K1C9_HUMAN]	0.96	1.06	1.10	1.17	1.15	1.19	1.15
P23528	Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3 - [COF1_HUMAN]	0.98	0.98	1.10	1.20	1.04	1.14	1.12
P00441	Superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD1 PE=1 SV=2 - [SODC_HUMAN]	1.02	1.02	1.10	1.02	1.07	1.00	1.05
P01861	Ig gamma-4 chain C region OS=Homo sapiens GN=IGHG4 PE=1 SV=1 - [IGHG4_HUMAN]	1.00	0.87	1.11	1.00	0.91	0.85	0.96
Q14103-4	Isoform 4 of Heterogeneous nuclear ribonucleoprotein D0 OS=Homo sapiens GN=HNRNPD - [HNRPD_HUMAN]	0.95	1.02	1.11	1.15	1.09	1.13	1.12
P00338	L-lactate dehydrogenase A chain OS=Homo sapiens GN=LDHA PE=1 SV=2 - [LDHA_HUMAN]	0.95	1.02	1.11	1.10	1.07	1.10	1.09
P61978-3	Isoform 3 of Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK - [HNRPK_HUMAN]	0.91	0.87	1.11	1.20	0.93	1.02	1.07
P0CG05	Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2 PE=1 SV=1 - [LAC2_HUMAN]	1.01	0.97	1.11	1.26	1.40	1.19	1.24
Q06830	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1 - [PRDX1_HUMAN]	1.03	0.96	1.12	1.05	1.05	1.04	1.06
P02749	Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1 SV=3 - [APOH_HUMAN]	1.08	0.97	1.12	1.02	0.98	0.90	1.01
Q8IVF2	Protein AHNAK2 OS=Homo sapiens GN=AHNAK2 PE=1 SV=2 - [AHNK2_HUMAN]	0.89	1.20	1.12	1.10	1.07	1.28	1.14
Q08188	Protein-glutamine gamma-glutamyltransferase E OS=Homo sapiens GN=TGM3 PE=1 SV=4 - [TGM3_HUMAN]	0.98	1.00	1.12	1.07	0.87	0.84	0.98
P20618	Proteasome subunit beta type-1 OS=Homo sapiens GN=PSMB1 PE=1 SV=2 - [PSB1_HUMAN]	1.08	1.06	1.13	1.03	1.10	1.02	1.07
P35321	Cornifin-A OS=Homo sapiens GN=SPRR1A PE=1 SV=2 - [SPR1A_HUMAN]	1.07	1.11	1.13	1.00	1.26	1.21	1.15
P01011	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2 - [AACT_HUMAN]	1.05	1.07	1.13	1.07	1.08	1.07	1.09
P60900	Proteasome subunit alpha type-6 OS=Homo sapiens GN=PSMA6 PE=1 SV=1 - [PSA6_HUMAN]	1.06	0.93	1.14	1.07	1.03	0.96	1.05
P40926	Malate dehydrogenase, mitochondrial OS=Homo sapiens GN=MDH2 PE=1 SV=3 - [MDHM_HUMAN]	0.97	0.93	1.15	1.21	1.02	1.07	1.11
Q05682-5	Isoform 5 of Caldesmon OS=Homo sapiens GN=CALD1 - [CALD1_HUMAN]	1.02	0.99	1.16	1.09	1.10	0.86	1.05
Q15582	Transforming growth factor-beta-induced protein ig-h3 OS=Homo sapiens GN=TGFBI PE=1 SV=1 - [BGH3_HUMAN]	0.91	1.01	1.16	1.09	1.03	1.13	1.11
P16070-15	Isoform 15 of CD44 antigen OS=Homo sapiens GN=CD44 - [CD44_HUMAN]	1.00	0.98	1.16	1.09	0.97	1.04	1.07
P06727	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3 - [APOA4_HUMAN]	1.01	0.96	1.17	1.20	1.08	1.12	1.14
P01620	Ig kappa chain V-III region SIE OS=Homo sapiens PE=1 SV=1 - [KV302_HUMAN]	0.97	0.97	1.17	1.19	1.10	1.12	1.15
P49721	Proteasome subunit beta type-2 OS=Homo sapiens GN=PSMB2 PE=1 SV=1 - [PSB2_HUMAN]	0.95	0.95	1.17	1.22	1.07	1.13	1.15
P04264	Keratin, type II cytoskeletal 1 OS=Homo sapiens GN=KRT1 PE=1 SV=6 - [K2C1_HUMAN]	1.04	0.99	1.17	1.09	1.14	1.03	1.11
P61586	Transforming protein RhoA OS=Homo sapiens GN=RHOA PE=1 SV=1 - [RHOA_HUMAN]	1.12	1.01	1.17	1.10	1.15	0.97	1.10
P08670	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4 - [VIME_HUMAN]	1.03	0.99	1.18	1.15	1.12	1.06	1.13
P01024	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2 - [CO3_HUMAN]	1.01	0.98	1.18	1.10	1.10	1.02	1.10
P60174-1	Isoform 2 of Triosephosphate isomerase OS=Homo sapiens GN=TPI1 - [TPIS_HUMAN]	1.07	0.99	1.19	1.10	1.17	1.05	1.13
P07585	Decorin OS=Homo sapiens GN=DCN PE=1 SV=1 - [PGS2_HUMAN]	0.91	0.99	1.19	1.11	1.30	1.50	1.27
P08107	Heat shock 70 kDa protein 1A/1B OS=Homo sapiens GN=HSPA1A PE=1 SV=5 - [HSP71_HUMAN]	1.07	0.99	1.19	1.10	1.08	1.03	1.10
P17174	Aspartate aminotransferase, cytoplasmic OS=Homo sapiens GN=GOT1 PE=1 SV=3 - [AATC_HUMAN]	0.83	1.24	1.20	0.87	1.38	1.08	1.13

P04406	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3 - [G3P_HUMAN]	0.97	0.96	1.21	1.25	1.18	1.20	1.21
P00491	Purine nucleoside phosphorylase OS=Homo sapiens GN=PNP PE=1 SV=2 - [PNPH_HUMAN]	0.97	0.89	1.22	1.24	1.11	1.13	1.17
Q15149	Plectin OS=Homo sapiens GN=PLEC PE=1 SV=3 - [PLEC_HUMAN]	0.86	0.97	1.22	1.30	1.14	1.22	1.22
P35579	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4 - [MYH9_HUMAN]	0.98	0.97	1.22	1.37	1.13	1.29	1.25
P02652	Apolipoprotein A-II OS=Homo sapiens GN=APOA2 PE=1 SV=1 - [APOA2_HUMAN]	0.95	0.97	1.23	1.38	1.04	1.27	1.23
P25786	Proteasome subunit alpha type-1 OS=Homo sapiens GN=PSMA1 PE=1 SV=1 - [PSA1_HUMAN]	1.01	0.98	1.23	1.21	1.13	1.10	1.17
P12277	Creatine kinase B-type OS=Homo sapiens GN=CKB PE=1 SV=1 - [KCRB_HUMAN]	0.91	1.10	1.23	1.34	1.39	**1.51**	1.37
P36952	Serpin B5 OS=Homo sapiens GN=SERPINB5 PE=1 SV=2 - [SPB5_HUMAN]	1.08	1.00	1.23	1.13	1.27	1.17	1.20
P07951-2	Isoform 2 of Tropomyosin beta chain OS=Homo sapiens GN=TPM2 - [TPM2_HUMAN]	0.96	0.98	1.25	1.40	**1.50**	**1.57**	1.43
P13647	Keratin, type II cytoskeletal 5 OS=Homo sapiens GN=KRT5 PE=1 SV=3 - [K2C5_HUMAN]	0.98	0.97	1.25	1.30	1.31	1.33	1.30
P28072	Proteasome subunit beta type-6 OS=Homo sapiens GN=PSMB6 PE=1 SV=4 - [PSB6_HUMAN]	1.03	0.96	1.26	1.21	1.18	1.13	1.20
P00558	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3 - [PGK1_HUMAN]	0.97	1.01	1.28	1.37	1.29	1.35	1.32
P00450	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1 - [CERU_HUMAN]	1.00	0.94	1.28	1.23	1.16	1.15	1.21
P61626	Lysozyme C OS=Homo sapiens GN=LYZ PE=1 SV=1 - [LYSC_HUMAN]	0.96	0.88	1.29	**1.61**	1.32	1.37	1.40
P07737	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2 - [PROF1_HUMAN]	0.99	0.93	1.29	1.34	1.18	1.23	1.26
P02792	Ferritin light chain OS=Homo sapiens GN=FTL PE=1 SV=2 - [FRIL_HUMAN]	1.06	0.98	1.30	1.19	1.22	1.13	1.21
Q9UBH0	Interleukin-36 receptor antagonist protein OS=Homo sapiens GN=IL36RN PE=1 SV=1 - [I36RA_HUMAN]	0.96	0.97	1.31	1.37	1.20	1.24	1.28
P26447	Protein S100-A4 OS=Homo sapiens GN=S100A4 PE=1 SV=1 - [S10A4_HUMAN]	0.91	0.95	1.31	1.47	1.18	1.35	1.33
P04433	Ig kappa chain V-III region VG (Fragment) OS=Homo sapiens PE=1 SV=1 - [KV309_HUMAN]	0.94	0.97	1.31	1.38	1.22	1.29	1.30
P06744	Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4 - [G6PI_HUMAN]	1.05	1.07	1.32	1.25	1.43	1.32	1.33
Q02487-2	Isoform 2B of Desmocollin-2 OS=Homo sapiens GN=DSC2 - [DSC2_HUMAN]	1.00	0.99	1.32	1.30	1.28	1.30	1.30
P06733	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2 - [ENOA_HUMAN]	1.06	0.96	1.33	1.24	1.23	1.16	1.24
P04040	Catalase OS=Homo sapiens GN=CAT PE=1 SV=3 - [CATA_HUMAN]	0.99	1.03	1.33	1.29	1.26	1.22	1.28
P18669	Phosphoglycerate mutase 1 OS=Homo sapiens GN=PGAM1 PE=1 SV=2 - [PGAM1_HUMAN]	1.04	1.02	1.34	1.28	1.32	1.27	1.30
P06703	Protein S100-A6 OS=Homo sapiens GN=S100A6 PE=1 SV=1 - [S10A6_HUMAN]	1.06	0.99	1.34	1.24	1.26	1.17	1.25
P68871	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2 - [HBB_HUMAN]	1.05	0.99	1.34	1.34	1.31	1.33	1.33
P30086	Phosphatidylethanolamine-binding protein 1 OS=Homo sapiens GN=PEBP1 PE=1 SV=3 - [PEBP1_HUMAN]	1.00	0.94	1.35	1.37	1.19	1.13	1.26
P01617	Ig kappa chain V-II region TEW OS=Homo sapiens PE=1 SV=1 - [KV204_HUMAN]	0.99	0.86	1.35	1.31	1.26	1.09	1.25
P29401	Transketolase OS=Homo sapiens GN=TKT PE=1 SV=3 - [TKT_HUMAN]	0.94	1.00	1.36	1.41	1.34	1.40	1.38
P07339	Cathepsin D OS=Homo sapiens GN=CTSD PE=1 SV=1 - [CATD_HUMAN]	1.04	1.00	1.36	1.28	1.30	1.25	1.30
P14618	Pyruvate kinase isozymes M1/M2 OS=Homo sapiens GN=PKM2 PE=1 SV=4 - [KPYM_HUMAN]	0.97	1.00	1.37	1.34	1.25	1.33	1.32
P52565	Rho GDP-dissociation inhibitor 1 OS=Homo sapiens GN=ARHGDIA PE=1 SV=3 - [GDIR1_HUMAN]	1.07	1.02	1.37	1.27	1.35	1.25	1.31
P02794	Ferritin heavy chain OS=Homo sapiens GN=FTH1 PE=1 SV=2 - [FRIH_HUMAN]	0.95	1.01	1.38	**1.58**	1.29	1.49	1.44
P02765	Alpha-2-HS-glycoprotein OS=Homo sapiens GN=AHSG PE=1 SV=1 - [FETUA_HUMAN]	0.89	0.88	1.39	**1.60**	1.22	1.33	1.39
P02647	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1 - [APOA1_HUMAN]	1.06	0.97	1.41	1.32	1.30	1.23	1.32
P02790	Hemopexin OS=Homo sapiens GN=HPX PE=1 SV=2 - [HEMO_HUMAN]	1.00	1.00	1.42	1.39	1.36	1.31	1.37
P30044-2	Isoform Cytoplasmic+peroxisomal of Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5 - [PRDX5_HUMAN]	1.13	1.00	1.43	1.33	1.36	1.20	1.33
P07602	Proactivator polypeptide OS=Homo sapiens GN=PSAP PE=1 SV=2 - [SAP_HUMAN]	1.15	1.13	1.44	1.10	1.46	1.15	1.29
P02751-10	Isoform 10 of Fibronectin OS=Homo sapiens GN=FN1 - [FINC_HUMAN]	1.01	0.97	1.44	**1.51**	1.49	**1.52**	1.49
P21333-2	Isoform 2 of Filamin-A OS=Homo sapiens GN=FLNA - [FLNA_HUMAN]	0.99	0.99	1.45	**1.66**	**1.57**	**1.56**	1.56
P25787	Proteasome subunit alpha type-2 OS=Homo sapiens GN=PSMA2 PE=1 SV=2 - [PSA2_HUMAN]	1.04	1.04	1.46	1.33	0.94	0.90	1.16
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1 - [AMBP_HUMAN]	0.96	1.00	1.46	1.45	1.39	1.35	1.41
P0C0L4	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=1 - [CO4A_HUMAN]	0.97	1.00	1.47	1.46	1.31	1.31	1.39
P04217	Alpha-1B-glycoprotein OS=Homo sapiens GN=A1BG PE=1 SV=4 - [A1BG_HUMAN]	0.97	1.02	1.47	1.49	1.42	1.32	1.42
B9A064	Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2 - [IGLL5_HUMAN]	0.91	0.95	1.48	**1.55**	1.37	**1.51**	1.48
P25789	Proteasome subunit alpha type-4 OS=Homo sapiens GN=PSMA4 PE=1 SV=1 - [PSA4_HUMAN]	0.94	1.03	1.48	**1.57**	1.42	**1.59**	1.52
P32119	Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5 - [PRDX2_HUMAN]	1.01	0.98	1.49	1.48	1.40	1.45	1.45
P02042	Hemoglobin subunit delta OS=Homo sapiens GN=HBD PE=1 SV=2 - [HBD_HUMAN]	1.02	0.99	1.49	1.32	1.37	1.26	1.36

P02774	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1 - [VTDB_HUMAN]	0.94	0.93	1.49	**1.60**	1.38	1.45	1.48
Q01518	Adenylyl cyclase-associated protein 1 OS=Homo sapiens GN=CAP1 PE=1 SV=5 - [CAP1_HUMAN]	1.07	0.99	1.49	1.34	1.48	1.25	1.39
P08603	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4 - [CFAH_HUMAN]	0.97	1.02	**1.51**	1.48	1.48	1.46	1.48
P01871	Ig mu chain C region OS=Homo sapiens GN=IGHM PE=1 SV=3 - [IGHM_HUMAN]	0.94	1.03	**1.51**	1.39	1.48	1.43	1.45
P01857	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [IGHG1_HUMAN]	1.05	0.90	**1.52**	1.37	1.35	1.21	1.36
P01860	Ig gamma-3 chain C region OS=Homo sapiens GN=IGHG3 PE=1 SV=2 - [IGHG3_HUMAN]	0.94	0.96	**1.55**	**1.64**	1.49	**1.52**	1.55
P80511	Protein S100-A12 OS=Homo sapiens GN=S100A12 PE=1 SV=2 - [S10AC_HUMAN]	0.98	0.96	**1.56**	**1.56**	1.44	1.45	1.50
P02533	Keratin, type I cytoskeletal 14 OS=Homo sapiens GN=KRT14 PE=1 SV=4 - [K1C14_HUMAN]	1.05	0.99	**1.59**	1.48	1.50	1.42	1.50
P08779	Keratin, type I cytoskeletal 16 OS=Homo sapiens GN=KRT16 PE=1 SV=4 - [K1C16_HUMAN]	0.98	0.97	**1.62**	**2.02**	1.43	**1.79**	1.71
P01008	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1 - [ANT3_HUMAN]	0.83	1.00	**1.63**	**1.93**	**1.57**	**1.87**	1.75
P02768	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2 - [ALBU_HUMAN]	1.04	0.96	**1.63**	**1.54**	**1.53**	1.48	1.55
Q6NXT2	Histone H3.3C OS=Homo sapiens GN=H3F3C PE=1 SV=3 - [H3C_HUMAN]	1.00	1.10	**1.64**	**1.63**	**1.64**	**1.60**	1.63
P02787	Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3 - [TRFE_HUMAN]	1.04	0.95	**1.65**	**1.54**	**1.55**	1.46	1.55
P01834	Ig kappa chain C region OS=Homo sapiens GN=IGKC PE=1 SV=1 - [IGKC_HUMAN]	1.09	0.91	**1.66**	1.37	1.44	1.29	1.44
P01859	Ig gamma-2 chain C region OS=Homo sapiens GN=IGHG2 PE=1 SV=2 - [IGHG2_HUMAN]	1.00	0.99	**1.67**	**1.75**	**1.51**	**1.59**	1.63
P13796	Plastin-2 OS=Homo sapiens GN=LCP1 PE=1 SV=6 - [PLSL_HUMAN]	0.96	0.94	**1.69**	**1.74**	**1.53**	**1.58**	1.63
P02763	Alpha-1-acid glycoprotein 1 OS=Homo sapiens GN=ORM1 PE=1 SV=1 - [A1AG1_HUMAN]	1.10	0.95	**1.69**	**1.53**	**1.55**	1.33	1.53
P00738	Haptoglobin OS=Homo sapiens GN=HP PE=1 SV=1 - [HPT_HUMAN]	1.07	0.97	**1.69**	**1.53**	**1.61**	1.48	1.58
P01876	Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1 PE=1 SV=2 - [IGHA1_HUMAN]	1.01	1.00	**1.71**	**1.67**	**1.67**	**1.62**	1.67
P09493-6	Isoform 6 of Tropomyosin alpha-1 chain OS=Homo sapiens GN=TPM1 - [TPM1_HUMAN]	1.02	0.95	**1.74**	**1.69**	1.40	1.36	1.55
P00915	Carbonic anhydrase 1 OS=Homo sapiens GN=CA1 PE=1 SV=2 - [CAH1_HUMAN]	1.03	0.91	**1.77**	**1.65**	1.45	1.43	1.57
P04196	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1 - [HRG_HUMAN]	1.06	0.91	**1.79**	**1.63**	**1.65**	1.44	1.63
P01009	Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1 SV=3 - [A1AT_HUMAN]	0.98	0.96	**1.79**	**1.78**	**1.81**	**1.82**	1.80
P69905	Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2 - [HBA_HUMAN]	1.07	0.99	**1.85**	**1.66**	**1.74**	**1.59**	1.71
Q04695	Keratin, type I cytoskeletal 17 OS=Homo sapiens GN=KRT17 PE=1 SV=2 - [K1C17_HUMAN]	0.98	0.94	**1.87**	**2.42**	**1.65**	**1.92**	1.97
P01023	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3 - [A2MG_HUMAN]	1.05	0.99	**1.90**	**1.83**	**1.66**	1.36	1.69
O60814	Histone H2B type 1-K OS=Homo sapiens GN=HIST1H2BK PE=1 SV=3 - [H2B1K_HUMAN]	1.05	0.95	**1.92**	**1.74**	**1.75**	**1.51**	1.73
P04259	Keratin, type II cytoskeletal 6B OS=Homo sapiens GN=KRT6B PE=1 SV=5 - [K2C6B_HUMAN]	0.97	0.91	**1.96**	**1.99**	**1.72**	**1.75**	1.85
P05164-2	Isoform H14 of Myeloperoxidase OS=Homo sapiens GN=MPO - [PERM_HUMAN]	0.94	1.00	**2.00**	**2.27**	**1.91**	**2.17**	2.09
Q96KK5	Histone H2A type 1-H OS=Homo sapiens GN=HIST1H2AH PE=1 SV=3 - [H2A1H_HUMAN]	1.02	0.98	**2.05**	**1.93**	**1.93**	**1.87**	1.94
O43707	Alpha-actinin-4 OS=Homo sapiens GN=ACTN4 PE=1 SV=2 - [ACTN4_HUMAN]	1.07	0.97	**2.06**	**1.89**	**1.89**	**1.74**	1.89
P02788	Lactotransferrin OS=Homo sapiens GN=LTF PE=1 SV=6 - [TRFL_HUMAN]	1.00	0.96	**2.12**	**2.06**	**1.93**	**1.94**	2.01
P80188	Neutrophil gelatinase-associated lipocalin OS=Homo sapiens GN=LCN2 PE=1 SV=2 - [NGAL_HUMAN]	1.01	0.93	**2.22**	**2.17**	**1.98**	**2.02**	2.10
P04179	Superoxide dismutase [Mn], mitochondrial OS=Homo sapiens GN=SOD2 PE=1 SV=2 - [SODM_HUMAN]	0.91	0.98	**2.29**	**2.43**	**2.19**	**2.38**	2.32
Q14624-2	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 - [ITIH4_HUMAN]	0.95	1.08	**2.39**	**2.49**	**2.50**	**2.61**	2.50
P01042-2	Isoform LMW of Kininogen-1 OS=Homo sapiens GN=KNG1 - [KNG1_HUMAN]	0.95	1.03	**2.52**	**2.44**	**2.38**	**2.17**	2.38
P02743	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2 - [SAMP_HUMAN]	1.01	0.97	**2.63**	**2.69**	**2.45**	**2.42**	2.55
P02538	Keratin, type II cytoskeletal 6A OS=Homo sapiens GN=KRT6A PE=1 SV=3 - [K2C6A_HUMAN]	1.02	0.94	**2.63**	**2.54**	**2.33**	**2.27**	2.44
P04004	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1 - [VTNC_HUMAN]	0.89	0.95	**2.71**	**2.74**	**2.52**	**2.71**	2.67
P08246	Neutrophil elastase OS=Homo sapiens GN=ELANE PE=1 SV=1 - [ELNE_HUMAN]	0.86	1.04	**2.79**	**2.81**	**2.79**	**2.81**	2.80
P10909-4	Isoform 4 of Clusterin OS=Homo sapiens GN=CLU - [CLUS_HUMAN]	0.93	0.96	**2.80**	**2.28**	**2.34**	**2.51**	2.48
P05155	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2 - [IC1_HUMAN]	0.91	0.94	**3.19**	**3.45**	**2.90**	**3.15**	3.17
P24158	Myeloblastin OS=Homo sapiens GN=PRTN3 PE=1 SV=3 - [PRTN3_HUMAN]	0.98	0.94	**3.45**	**3.48**	**3.13**	**3.18**	3.31
P02671-2	Isoform 2 of Fibrinogen alpha chain OS=Homo sapiens GN=FGA - [FIBA_HUMAN]	0.99	0.96	**3.60**	**3.57**	**3.21**	**3.33**	3.43
P62805	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2 - [H4_HUMAN]	0.98	0.95	**3.66**	**3.41**	**3.28**	**2.99**	3.34
P02679-2	Isoform Gamma-A of Fibrinogen gamma chain OS=Homo sapiens GN=FGG - [FIBG_HUMAN]	1.00	0.96	**4.58**	**4.50**	**4.27**	**3.93**	4.32
P02675	Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1 SV=2 - [FIBB_HUMAN]	1.04	0.96	**4.78**	**4.89**	**3.98**	**4.35**	4.50