Supplementary Table. Protein names, accession numbers and spectral counts of those proteins identified in this study S: semi-tryptic; TCA: trichloroacetic acid; UC: ultracentrifugation

Protein name	UniProt AC	UniProt ID	Average spectral counts				Spectral Counts																					
			TCA	TCA-S	UC	UC-S	TCA1	TCA2	TCA3	TCA4	TCA5	TCA6	TCA-S1	TCA-S2	TCA-S3	TCA-S4	TCA-S5	TCA-S6	UC1	UC2	UC3	UC4	UC5	UC6	UC-S1	UC-S2	UC-S3	UC-S4	UC-S5	UC-S6
Acid sphingomyelinase-like phosphodiesterase 3b	Q92485	ASM3B_HUMAN	0	0	4	4	0	0	0	0	0	0	0	0	0	0	0	0	10	5	5	0	5	0	8	5	5	0	4	0
Alkaline phosphatase, tissue-nonspecific isozyme	P05186	PPBT_HUMAN	0	0	2	2	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	11	0	0	0	0	0	9
Annexin A11	P50995	ANX11_HUMAN	1	0	1	2	0	0	0	0	5	0	0	0	0	0	2	0	0	0	0	0	5	3	4	0	0	0	4	3
Calmodulin-like protein 3	P27482	CALL3_HUMAN	0	0	1	1	0	0	0	0	1	1	0	0	0	0	1	1	0	0	0	7	0	0	0	0	0	5	0	0
Cathepsin G	P08311	CATG_HUMAN	0	1	6	4	0	0	0	0	0	2	0	0	0	0	0	3	10	0	0	0	0	25	8	0	0	0	0	18
Complement component C9	P02748	CO9_HUMAN	0	1	1	1	0	0	0	0	0	0	1	1	1	1	1	1	0	0	0	0	0	8	0	0	0	0	0	6
C-reactive protein	P02741	CRP_HUMAN	0	0	2	1	0	0	0	0	0	0	0	0	0	0	0	0	10	0	0	0	0	0	8	0	0	0	0	0
Ectonucleotide pyrophosphatase/phosphodiesterase family member 7	Q6UWV6	ENPP7_HUMAN	0	0	3	3	2	0	0	0	0	0	2	0	0	0	0	0	5	5	0	0	5	3	4	5	0	0	4	3
Eosinophil peroxidase	P11678	PERE_HUMAN	0	0	9	8	0	1	0	0	0	1	0	1	0	0	0	1	34	0	0	0	15	3	29	0	0	0	16	3
Ferritin heavy chain	P02794	FRIH_HUMAN	0	0	6	5	1	0	1	0	0	0	0	0	1	0	0	0	5	27	5	0	0	0	0	22	5	0	0	0
Histone H2B type 1-K	O60814	H2B1K_HUMAN	0	0	1	2	0	0	0	0	2	0	0	0	0	0	1	0	0	5	0	0	0	0	0	9	0	0	0	0
Isoform 1 of Plakophilin-1	Q13835-2	PKP1_HUMAN	0	0	1	1	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	5	0	0	0	0	0	4	0
Mucin-1	P15941	MUC1_HUMAN	0	0	5	4	0	0	0	0	0	1	0	0	0	0	0	2	0	0	18	0	0	11	0	0	16	0	0	9
Pyruvate kinase isozymes M1/M2	P14618	KPYM_HUMAN	0	1	1	1	0	0	0	0	2	0	0	1	0	1	5	0	0	0	0	7	0	0	0	0	0	5	0	0
WD repeat-containing protein 67	Q96DN5	WDR67_HUMAN	0	0	0	2	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	9	5	0	0	0
Zinc finger protein 337	Q9Y3M9	ZN337_HUMAN	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	12	0	0
10 kDa heat shock protein, mitochondrial	P61604	CH10_HUMAN	3	3	0	0	1	1	1	5	5	2	1	1	1	5	5	2	0	0	0	0	0	0	0	0	0	0	0	0
14-3-3 protein epsilon	P62258	1433E_HUMAN	3	2	1	1	0	1	9	0	5	0	0	1	7	0	5	0	0	0	0	7	0	0	0	0	0	5	0	0
14-3-3 protein gamma	P61981	1433G_HUMAN	2	1	1	1	0	1	2	0	6	0	0	1	2	0	5	0	0	0	0	7	0	0	0	0	0	5	0	0
14-3-3 protein zeta/delta	P63104	1433Z_HUMAN	4	4	2	2	0	3	11	0	12	0	0	3	11	0	10	0	0	0	0	14	0	0	0	0	0	12	0	0
2,4-dienoyl-CoA reductase, mitochondrial	Q16698	DECR_HUMAN	3	3	0	0	0	5	7	0	4	0	0	6	8	0	5	0	0	0	0	0	0	0	0	0	0	0	0	0
2-oxoglutarate dehydrogenase, mitochondrial	Q02218	ODO1_HUMAN	1	1	0	0	0	1	2	0	0	0	0	1	2	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
3-hydroxybutyrate dehydrogenase type 2	Q9BUT1	BDH2_HUMAN	2	2	0	0	2	2	3	1	6	0	2	2	3	1	5	0	0	0	0	0	0	0	0	0	0	0	0	0
3-ketoacyl-CoA thiolase, mitochondrial	P42765	THIM_HUMAN	2	3	0	0	0	2	10	0	2	0	0	3	10	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0
40S ribosomal protein S15a	P62244	RS15A_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
40S ribosomal protein S7	P62081	RS7_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
4-aminobutyrate aminotransferase, mitochondrial	P80404	GABT_HUMAN	1	1	0	0	1	1	2	1	2	0	1	1	2	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
55 kDa erythrocyte membrane protein	Q00013	EM55_HUMAN	1	1	0	0	0	1	2	0	0	0	0	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
5'-nucleotidase	P21589	5NTD_HUMAN	1	1	0	0	0	0	0	0	0	8	0	0	0	0	0	8	0	0	0	0	0	0	0	0	0	0	0	0
60S acidic ribosomal protein P0	P05388	RLA0_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
60S acidic ribosomal protein P2	P05387	RLA2_HUMAN	1	1	0	0	0	1	2	1	1	1	0	2	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
6-phosphogluconate dehydrogenase, decarboxylating	P52209	6PGD_HUMAN	1	1	0	0	0	0	4	0	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
78 kDa glucose-regulated protein	P11021	GRP78_HUMAN	7	6	1	0	2	7	15	5	8	3	2	4	13	5	7	3	0	0	0	7	0	0	0	0	0	0	0	0
Acetyl-CoA acetyltransferase, mitochondrial	P24752	THIL_HUMAN	1	1	0	0	0	0	4	0	2	0	0	0	4	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Acid ceramidase	Q13510	ASAH1_HUMAN	0	0	0	0	0	0	0	0	0	2	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Aconitate hydratase, mitochondrial	Q99798	ACON_HUMAN	2	2	0	0	0	0	10	0	2	0	0	0	8	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Actin, cytoplasmic 1	P60709	ACTB_HUMAN	15	14	4	3	1	11	21	11	26	17	1	9	21	8	26	17	0	0	0	21	0	0	0	0	0	17	0	0
Actin-related protein 2/3 complex subunit 4	P59998	ARPC4_HUMAN	0	0	0	0	0	0	1	0	1	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Actin-related protein 3	P61158	ARP3_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Adenosine deaminase	P00813	ADA_HUMAN	12	11	0	0	18	8	10	16	15	7	17	5	9	14	13	5	0	0	0	0	0	0	0	0	0	0	0	0
Adenylyl cyclase-associated protein 1	Q01518	CAP1_HUMAN	5	4	0	0	8	4	4	5	5	2	6	4	2	5	5	2	0	0	0	0	0	0	0	0	0	0	0	0
Adseverin	Q9Y6U3	ADSV_HUMAN	1	1	0	0	0	1	0	1	1	1	0	1	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Afamin	P43652	AFAM_HUMAN	1	1	0	0	0	1	6	0	1	0	0	0	4	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Aflatoxin B1 aldehyde reductase member 3	O95154	ARK73_HUMAN	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Alcohol dehydrogenase [NADP+]	P14550	AK1A1_HUMAN	8	9	0	0	6	8	9	14	12	0	5	12	8	17	10	0	0	0	0	0	0	0	0	0	0	0	0	0
Alcohol dehydrogenase 1C	P00326	ADH1G_HUMAN	7	7	0	0	11	8	10	2	12	0	10	8	9	2	12	0	0	0	0	0	0	0	0	0	0	0	0	0
Aldehyde dehydrogenase, dimeric NADP-preferring	P30838	AL3A1_HUMAN	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Aldehyde dehydrogenase, mitochondrial	P05091	ALDH2_HUMAN	1	1	1	1	1	1	2	0	2	0	1	1	2	0	2	0	0	0	0	0	5	3	0	0	0	0	4	3
Aldo-keto reductase family 1 member B10	O60218	AK1BA_HUMAN	24	25	0	0	25	22	25	26	28	18	24	25	26	25	30	18	0	0	0	0	0	0	0	0	0	0	0	0
Aldo-keto reductase family 1 member C1	Q04828	AK1C1_HUMAN	4	6	0	0	4	3	4	4	5	1	6	6	7	5	7	2	0	0	0	0	0	0	0	0	0	0	0	0
Aldo-keto reductase family 1 member C3	P42330	AK1C3_HUMAN	5	6	0	0	5	8	6	4	6	1	5	10	7	5	7	2	0	0	0	0	0	0	0	0	0	0	0	0
Aldose 1-epimerase	Q96C23	GALM_HUMAN	3	2	1	1	4	3	2	4	2	1	3	3	2	4	1	1	0	0	0	0	5	0	0	0	0	0	4	0
Alpha-1-acid glycoprotein 1	P02763	A1AG1_HUMAN	23	22	18	16	14	15	17	20	27	44	12	15	16	18	27	41	5	5	30	0	24	46	8	5	21	0	21	39
Alpha-1-acid glycoprotein 2	P19652	A1AG2_HUMAN	9	9	11	10	4	10	10	6	9	17	3	9	8	6	8	18	5	5	24	0	20	14	8	5	16	0	16	15
Alpha-1-antichymotrypsin	P01011	AACT_HUMAN	13	12	1	1	7	11	10	17	9	22	8	10	9	15	9	19	0	0	0	0	0	3	0	0	0	0	0	3
Alpha-1-antitrypsin	P01009	A1AT_HUMAN	731	708	659	707	91	156	138	116	158	72	84	158	133	112	152	69	110	78	114	183	131	43	119	90	142	182	129	45
Alpha-1B-glycoprotein	P04217	A1BG_HUMAN	10	10	0	0	6	12	12	11	16	3	6	12	11	9	17	3	0	0	0	0	0	0	0	0	0	0	0	0
Alpha-2-antiplasmin	P08697	A2AP_HUMAN	1	1	0	0	0	2	1	1	1	0	0	3	2	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Alpha-2-HS-glycoprotein	P02765	FETUA_HUMAN	1	0	0	0	0	0	0	0	1	2	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Alpha-2-macroglobulin	P01023	A2MG_HUMAN	122	114	212	189	35	216	148	141	142	51	29	204	142	128	135	44	332	83	257	37	357	207	293	72	242	21	318	189
Alpha-2-macroglobulin-like protein 1	A8K2U0	A2ML1_HUMAN	1	3	0	0	0	0	0	0	0	7	1	2	1	1	1	10	0	0	0	0	0	0	0	0	0	0	0	0
Alpha-actinin-4	O43707	ACTN4_HUMAN	12	12	0	0	2	12	19	8	26	7	2	11	17	9	23	7	0	0	0	0	0	0	0	0	0	0	0	0
Alpha-amylase 1	P04745	AMY1_HUMAN	166	172	322	356	176	157	140	96	106	322	182	159	144	104	113	330	254	165	503	337	227	447	293	179	478	401	270	515
Alpha-amylase 2B	P19961	AMY2B_HUMAN	166	175	381	416	191	168	150	108	114	262	202	176	160	121	127	266	332	225	598	405	309	418	368	243	578	472	362	473
Alpha-enolase	P06733	ENOA_HUMAN	4	4	4	3	0	0	13	2	6	3	0	0	11	2	5	3	0	0	0	21	0	0	0	0	0	17	0	0
Aminoacylase-1	Q03154	ACY1_HUMAN	10	9	0	0	12	10	10	10	12	5	11	9	9	8	12	3	0	0	0	0	0	0	0	0	0	0	0	0
Aminopeptidase N	P15144	AMPN_HUMAN	51	48	31	27	64	21	20	65	62	73	61	22	17	63	58	69	15	16	12	7	53	82	12	13	11	5	44	75
Angiotensin-converting enzyme 2	Q9BYF1	ACE2_HUMAN	2	2	0	0	2	0	0	4	1	5	1	0	0	4	1	5	0	0	0	0	0	0	0	0	0	0	0	0
Angiotensinogen	P01019	ANGT_HUMAN	1	2	0	0	0	2	6	0	0	0	0	2	7	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Annexin A1	P04083	ANXA1_HUMAN	1	1	1	1	0	0	0	0	5	1	0	0	0	0	5	1	0	0	0	7	0	0	0	0	0	5	0	0
Annexin A10	Q9UJ72	ANX10_HUMAN	0	0	0	0	2	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Annexin A13	P27216	ANX13_HUMAN	2	2	0	0	0	0	0	0	13	0	0	0	0	0	13	0	0	0	0	0	0	0	0	0	0	0	0	0
Annexin A3	P12429	ANXA3_HUMAN	1	1	0	0	0	0	0	0	0	5	0	0	0	0	0	5	0	0	0	0	0	0	0	0	0	0	0	0
Annexin A4	P09525	ANXA4_HUMAN	4	4	7	6	0	2	4	4	15	1	0	1	4	5	14	1	10	0	5	0	15	11	8	0	5	0	12	9
Anterior gradient protein 2 homolog	O95994	AGR2_HUMAN	1	1	0	0	0	1	3	0	0	0	0	1	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0

Antileukoproteinase	P03973	SLPI_HUMAN	3	3	0	1	0	0	0	1	0	14	0	0	0	1	0	14	0	0	0	0	0	0	0	0	0	0	0	3
Antithrombin-III	P01008	ANT3_HUMAN	10	10	6	6	3	14	13	8	13	8	3	14	12	8	13	8	15	0	12	0	9	0	12	0	11	5	8	0
Apolipoprotein A-I	P02647	APOA1_HUMAN	2	2	0	0	0	0	10	0	0	0	0	0	13	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Apolipoprotein B-100	P04114	APOB_HUMAN	2	2	0	0	0	1	12	0	0	1	0	1	12	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Apolipoprotein C-III	P02656	APOC3_HUMAN	0	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Apolipoprotein D	P05090	APOD_HUMAN	3	2	0	0	2	7	6	1	0	3	1	5	4	1	0	2	0	0	0	0	0	0	0	0	0	0	0	0
Aromatic-L-amino-acid decarboxylase	P20711	DDC_HUMAN	1	1	0	0	0	1	1	0	1	0	0	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Aspartate aminotransferase, cytoplasmic	P17174	AATC_HUMAN	6	5	0	0	9	3	4	4	11	6	8	3	4	4	8	5	0	0	0	0	0	0	0	0	0	0	0	0
Aspartate aminotransferase, mitochondrial	P00505	AATM_HUMAN	4	4	0	0	5	2	2	5	5	5	5	2	3	6	5	5	0	0	0	0	0	0	0	0	0	0	0	0
ATP synthase subunit alpha, mitochondrial	P25705	ATPA_HUMAN	0	0	0	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
ATP synthase subunit beta, mitochondrial	P06576	ATPB_HUMAN	1	1	0	0	0	0	0	0	4	0	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0
ATP synthase subunit g, mitochondrial	O75964	ATP5L_HUMAN	1	1	0	0	0	0	0	5	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Attractin	O75882	ATRN_HUMAN	3	2	0	0	2	1	1	6	4	2	1	1	1	4	2	2	0	0	0	0	0	0	0	0	0	0	0	0
Azurocidin	P20160	CAP7_HUMAN	1	1	1	1	0	0	0	0	0	8	0	0	0	0	0	8	0	0	0	0	0	8	0	0	0	0	0	6
Bactericidal/permeability-increasing protein-like 1	Q8N4F0	BPIL1_HUMAN	1	1	0	0	0	0	0	0	0	7	0	0	0	0	0	7	0	0	0	0	0	0	0	0	0	0	0	0
Beta-2-glycoprotein 1	P02749	APOH_HUMAN	8	8	0	0	2	5	6	8	9	15	2	5	7	8	9	14	0	0	0	0	0	0	0	0	0	0	0	0
Beta-2-microglobulin	P61769	B2MG_HUMAN	1	1	0	0	0	0	0	0	2	2	0	0	1	0	1	2	0	0	0	0	0	0	0	0	0	0	0	0
Beta-microseminoprotein	P08118	MSMB_HUMAN	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0
Bifunctional ATP-dependent dihydroxyacetone kinase/FAD-AMP lyase (cyclizing)	Q3LXA3	DHAK_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Bile salt-activated lipase	P19835	CEL_HUMAN	110	108	24	20	212	46	69	161	148	22	215	41	62	160	150	20	39	38	5	30	20	11	37	32	5	21	16	9
C4b-binding protein alpha chain	P04003	C4BPA_HUMAN	3	2	0	0	1	5	8	1	0	0	1	4	8	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Cadherin-1	P12830	CADH1_HUMAN	1	1	0	0	1	0	0	0	2	1	1	0	0	0	2	1	0	0	0	0	0	0	0	0	0	0	0	0
Cadherin-17	Q12864	CAD17_HUMAN	2	2	0	0	6	0	1	6	1	0	6	0	0	6	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Cadherin-related family member 2	Q9BYE9	CDHR2_HUMAN	20	19	4	4	25	2	1	25	24	44	26	1	1	24	23	40	0	5	0	0	9	11	0	5	0	0	8	9
Calcium-activated chloride channel regulator 1	A8K7I4	CLCA1_HUMAN	27	24	76	83	32	15	24	59	21	10	27	13	22	54	19	9	91	78	54	59	140	36	107	67	62	66	153	45
Calcium-binding protein p22	Q99653	CHP1_HUMAN	1	1	0	0	0	1	1	1	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Calmodulin	P62158	CALM_HUMAN	4	4	0	0	0	2	3	7	6	6	0	2	3	6	6	6	0	0	0	0	0	0	0	0	0	0	0	0
Calmodulin-like protein 5	Q9NZT1	CALL5_HUMAN	1	1	1	1	0	0	0	0	2	2	0	0	0	0	2	3	0	0	0	7	0	0	0	0	0	5	0	0
Calnexin	P27824	CALX_HUMAN	0	1	0	0	0	0	0	1	1	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Calpain-1 catalytic subunit	P07384	CAN1_HUMAN	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Calreticulin	P27797	CALR_HUMAN	5	4	0	0	6	5	4	6	7	1	5	5	2	6	6	1	0	0	0	0	0	0	0	0	0	0	0	0
Carbonic anhydrase 2	P00918	CAH2_HUMAN	10	10	4	4	9	12	15	13	5	6	10	12	13	12	6	7	0	16	5	0	0	3	0	13	5	0	0	3
Carbonic anhydrase 6	P23280	CAH6_HUMAN	2	5	0	0	0	0	0	0	0	14	0	0	0	0	0	28	0	0	0	0	0	0	0	0	0	0	0	0
Carbonyl reductase [NADPH] 1	P16152	CBR1_HUMAN	8	7	0	0	0	10	29	1	5	0	0	9	27	1	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Carboxymethylenebutenolidase homolog	Q96DG6	CMBL_HUMAN	1	1	0	0	1	2	1	0	1	0	1	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Carboxypeptidase A1	P15085	CBPA1_HUMAN	270	255	274	259	358	267	249	267	353	127	336	252	237	241	346	115	254	419	317	309	222	125	240	405	336	248	209	114
Carboxypeptidase A2	P48052	CBPA2_HUMAN	22	22	10	11	11	36	32	25	21	9	11	37	31	25	20	8	10	0	42	0	5	0	8	0	48	0	4	3
Carboxypeptidase B	P15086	CBPB1_HUMAN	196	194	170	164	357	170	159	187	239	65	368	153	151	186	238	67	135	298	216	89	188	93	148	301	215	54	181	87
Carcinoembryonic antigen-related cell adhesion molecule 5	P06731	CEAM5_HUMAN	2	3	6	5	2	0	0	2	2	7	2	0	0	4	2	8	5	5	0	0	15	8	4	5	0	0	12	6
Carnitine O-palmitoyltransferase 2, mitochondrial	P23786	CPT2_HUMAN	1	1	0	0	1	0	1	1	1	1	1	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Caspase-14	P31944	CASPE_HUMAN	1	1	0	0	0	0	0	0	2	1	0	0	0	0	2	1	0	0	0	0	0	0	0	0	0	0	0	0
Cathelicidin antimicrobial peptide	P49913	CAMP_HUMAN	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Cathepsin D	P07339	CATD_HUMAN	5	4	0	0	3	3	4	4	6	8	2	3	3	4	6	7	0	0	0	0	0	0	0	0	0	0	0	0
Cathepsin Z	Q9UBR2	CATZ_HUMAN	1	1	0	0	1	1	1	0	0	2	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
CD59 glycoprotein	P13987	CD59_HUMAN	1	1	0	0	0	0	0	0	0	3	0	0	0	0	0	5	0	0	0	0	0	0	0	0	0	0	0	0
Cell surface A33 antigen	Q99795	GPA33_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Ceruloplasmin	P00450	CERU_HUMAN	13	12	1	1	1	15	15	8	18	22	1	13	14	7	16	19	0	0	0	0	0	3	0	0	0	0	0	3
Chloride intracellular channel protein 1	O00299	CLIC1_HUMAN	1	1	0	0	0	1	6	0	1	0	0	2	5	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Choline transporter-like protein 4	Q53GD3	CTL4_HUMAN	1	1	3	2	1	0	0	1	0	1	1	0	0	1	0	1	0	0	0	7	9	3	0	0	0	0	8	3
Chymotrypsin-C	Q99895	CTRC_HUMAN	49	48	42	30	79	57	44	65	32	16	83	55	39	63	33	14	48	49	84	30	29	11	46	27	68	21	8	9
Chymotrypsin-like elastase family member 2A	P08217	CEL2A_HUMAN	159	162	65	58	210	160	160	189	154	79	229	172	166	182	153	69	71	94	90	66	53	14	66	104	89	38	40	12
Chymotrypsin-like elastase family member 2B	P08218	CEL2B_HUMAN	56	80	20	18	80	54	58	67	54	24	107	89	84	95	70	36	19	33	24	21	20	3	21	35	27	12	12	3
Chymotrypsin-like elastase family member 3A	P09093	CEL3A_HUMAN	95	101	151	148	134	91	116	124	84	23	143	96	126	130	85	25	130	187	306	162	120	3	140	166	315	144	120	3
Chymotrypsin-like elastase family member 3B	P08861	CEL3B_HUMAN	41	60	63	79	61	37	45	45	44	12	90	54	71	71	59	16	63	83	90	82	58	3	87	104	126	78	72	6
Chymotrypsin-like protease CTRL-1	P40313	CTRL_HUMAN	9	8	0	0	9	10	12	13	7	0	9	9	12	12	8	0	0	0	0	0	0	0	0	0	0	0	0	0
Chymotrypsinogen B2	Q6GPI1	CTRB2_HUMAN	136	140	14	21	226	140	122	149	127	50	221	148	130	155	135	48	15	49	5	0	5	8	21	59	16	12	4	15
Citrate synthase, mitochondrial	O75390	CISY_HUMAN	1	1	0	0	0	1	2	0	1	0	0	1	2	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Coatomer subunit delta	P48444	COPD_HUMAN	0	0	0	0	0	0	1	0	1	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Coatomer subunit gamma	Q9Y678	COPG_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Cochlin	O43405	COCH_HUMAN	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Cofilin-1	P23528	COF1_HUMAN	1	1	0	0	0	0	2	0	2	0	0	0	2	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Colipase	P04118	COL_HUMAN	12	16	5	6	5	19	17	20	9	0	4	21	22	31	14	1	0	11	12	0	9	0	0	9	11	5	8	0
Complement C1q subcomponent subunit B	P02746	C1QB_HUMAN	1	1	0	0	0	1	3	0	1	1	0	1	2	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Complement C1q subcomponent subunit C	P02747	C1QC_HUMAN	1	1	0	0	0	1	3	0	0	0	0	1	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Complement C2	P06681	CO2_HUMAN	0	1	0	0	0	0	0	0	0	0	3	1	1	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Complement C3	P01024	CO3_HUMAN	107	101	15	13	47	125	157	106	113	91	46	119	152	99	106	83	10	0	30	0	34	14	8	0	27	0	28	12
Complement C4-B	P0C0L5	CO4B_HUMAN	17	16	1	1	1	20	31	10	18	22	1	18	30	9	17	22	0	0	5	0	0	3	0	0	5	0	0	3
Complement C5	P01031	CO5_HUMAN	1	2	0	0	0	1	2	1	1	0	1	2	2	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Complement factor B	P00751	CFAB_HUMAN	3	3	0	0	0	2	9	0	5	1	0	3	7	0	5	1	0	0	0	0	0	0	0	0	0	0	0	0
Complement factor H	P08603	CFAH_HUMAN	4	4	0	0	1	2	13	0	7	3	0	3	11	0	6	3	0	0	0	0	0	0	0	0	0	0	0	0
Complement factor H-related protein 1	Q03591	FHR1_HUMAN	0	0	0	0	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0
Cornulin	Q9UBG3	CRNN_HUMAN	0	0	0	0	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0
Coronin-1B	Q9BR76	COR1B_HUMAN	0	0	0	0	0	0	1	0	1	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Corticosteroid-binding globulin	P08185	CBG_HUMAN	0	1	0	0	0	1	0	0	0	0	1	2	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
CUB and zona pellucida-like domain-containing protein 1	Q86UP6	CUZD1_HUMAN	14	15	97	130	57	10	7	6	5	0	62	10	7	5	5	0	115	209	48	59	102	50	140	337	48	66	133	54
Cystatin-A	P01040	CYTA_HUMAN	1	1	0	0	1	0	0	0	0	6	1	0	0	0	0	7	0	0	0	0	0	0	0	0	0	0	0	0
Cystatin-B	P04080	CYTB_HUMAN	3	2	0	0	1	0	1	2	6	6	1	0	0	2	6	5	0	0	0	0	0	0	0	0	0	0	0	0

Cystatin-S	P01036	CYTS_HUMAN	4	3	1	1	0	0	0	0	0	22	0	0	0	0	0	20	0	0	0	0	0	3	0	0	0	0	0	3
Cystatin-SA	P09228	CYTT_HUMAN	2	2	0	0	0	0	0	0	0	13	0	0	0	0	0	11	0	0	0	0	0	0	0	0	0	0	0	0
Cystatin-SN	P01037	CYTN_HUMAN	2	2	0	0	0	0	0	0	0	13	0	0	0	0	0	13	0	0	0	0	0	0	0	0	0	0	0	0
Cysteine-rich secretory protein 3	P54108	CRIS3_HUMAN	1	1	0	0	0	0	0	0	0	5	0	0	0	0	0	7	0	0	0	0	0	0	0	0	0	0	0	0
Cytochrome c	P99999	CYC_HUMAN	6	5	1	1	7	3	6	7	11	2	7	2	4	7	9	2	5	0	0	0	0	0	4	0	0	0	0	0
Cytochrome c oxidase subunit 2	P00403	COX2_HUMAN	2	2	0	0	1	0	1	6	1	1	2	0	1	7	2	2	0	0	0	0	0	0	0	0	0	0	0	0
Cytochrome c oxidase subunit 4 isoform 1, mitochondrial	P13073	COX41_HUMAN	1	1	0	0	0	0	0	2	2	2	0	0	0	2	2	2	0	0	0	0	0	0	0	0	0	0	0	0
Cytochrome c oxidase subunit 5A, mitochondrial	P20674	COX5A_HUMAN	1	2	0	0	0	0	0	1	4	1	0	0	0	4	5	2	0	0	0	0	0	0	0	0	0	0	0	0
Cytochrome c oxidase subunit 5B, mitochondrial	P10606	COX5B_HUMAN	0	0	0	0	0	0	0	0	1	1	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Cytochrome c oxidase subunit 6C	P09669	COX6C_HUMAN	1	1	0	0	0	1	0	2	2	1	0	1	0	2	3	1	0	0	0	0	0	0	0	0	0	0	0	0
Cytochrome c oxidase subunit 7A2, mitochondrial	P14406	CX7A2_HUMAN	2	2	0	0	2	1	2	2	5	1	2	1	2	2	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Cytoplasmic aconitate hydratase	P21399	ACOC_HUMAN	1	1	0	0	0	1	4	0	0	0	0	1	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Cytosolic non-specific dipeptidase	Q96KP4	CNDP2_HUMAN	4	4	1	1	0	8	6	1	6	1	0	8	7	1	6	2	0	0	0	0	0	5	0	0	0	0	4	0
D-dopachrome decarboxylase	P30046	DOPD_HUMAN	4	3	0	0	0	7	9	1	6	2	0	5	8	0	5	1	0	0	0	0	0	0	0	0	0	0	0	0
Defensin-5	Q01523	DEF5_HUMAN	6	5	0	0	3	2	2	4	8	16	3	2	1	4	7	15	0	0	0	0	0	0	0	0	0	0	0	0
Defensin-6	Q01524	DEF6_HUMAN	1	0	0	0	0	0	0	0	1	2	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Dehydrogenase/reductase SDR family member 11	Q6UWP2	DHR11_HUMAN	3	3	0	0	1	5	6	2	4	1	1	6	4	4	3	1	0	0	0	0	0	0	0	0	0	0	0	0
Deoxyribonuclease-1	P24855	DNAS1_HUMAN	3	3	0	0	3	0	1	6	2	5	2	0	1	6	2	5	0	0	0	0	0	0	0	0	0	0	0	0
Dermcidin	P81605	DCD_HUMAN	8	7	12	10	9	8	6	11	8	5	8	8	4	9	7	5	10	5	12	21	15	11	8	5	11	17	12	9
Desmoglein-1	Q02413	DSG1_HUMAN	1	1	5	5	0	0	1	1	2	2	0	0	1	0	2	1	0	0	5	14	9	0	0	0	5	17	8	0
Desmoplakin	P15924	DESP_HUMAN	5	5	4	3	0	1	0	1	20	6	0	1	1	1	20	7	0	0	0	7	15	3	0	0	0	5	12	3
Dihydrolipoyl dehydrogenase, mitochondrial	P09622	DLDH_HUMAN	2	2	0	0	2	1	3	1	2	0	2	1	3	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Dihydrolipoyllysine-residue succinyltransferase, 2-oxoglutarate dehydrogenase	P36957	ODO2_HUMAN	2	2	0	0	2	1	1	2	2	1	2	1	1	2	2	1	0	0	0	0	0	0	0	0	0	0	0	0
Dipeptidyl peptidase 4	P27487	DPP4_HUMAN	4	3	4	3	7	1	2	4	5	5	6	1	1	2	3	5	0	0	0	0	9	14	0	0	0	0	8	12
DNA-(apurinic or apyrimidinic site) lyase	P27695	APEX1_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit	P39656	OST48_HUMAN	0	0	0	0	0	1	0	1	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Dynein heavy chain 8, axonemal	Q96JB1	DYH8_HUMAN	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Echinoderm microtubule-associated protein-like 4	Q9HC35	EMAL4_HUMAN	0	0	0	0	0	0	1	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Elongation factor 1-gamma	P26641	EF1G_HUMAN	1	1	0	0	3	1	0	0	1	0	2	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Elongation factor 2	P13639	EF2_HUMAN	2	3	0	0	0	2	4	0	6	1	0	3	4	1	6	2	0	0	0	0	0	0	0	0	0	0	0	0
Enteropeptidase	P98073	ENTK_HUMAN	27	26	42	37	23	20	13	39	32	35	20	18	13	39	34	33	15	38	0	110	49	39	12	32	0	99	40	37
Eosinophil cationic protein	P12724	ECP_HUMAN	1	1	0	0	0	1	0	0	1	2	1	1	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Epididymal secretory protein E1	P61916	NPC2_HUMAN	1	1	0	0	0	1	0	0	1	2	0	1	0	0	1	2	0	0	0	0	0	0	0	0	0	0	0	0
Epithelial cell adhesion molecule	P16422	EPCAM_HUMAN	2	2	0	0	2	1	1	4	2	2	2	1	1	4	2	2	0	0	0	0	0	0	0	0	0	0	0	0
Epoxide hydrolase 1	P07099	HYEP_HUMAN	2	2	0	0	1	0	1	5	2	1	1	0	1	4	2	1	0	0	0	0	0	0	0	0	0	0	0	0
Eukaryotic initiation factor 4A-I	P60842	IF4A1_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Ezrin	P15311	EZRI_HUMAN	1	1	0	0	0	1	3	1	2	0	0	1	2	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Fatty acid-binding protein, intestinal	P12104	FABPI_HUMAN	3	3	0	0	0	8	10	0	0	0	0	8	9	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Fatty acid-binding protein, liver	P07148	FABPL_HUMAN	15	15	0	0	1	20	18	19	20	14	1	21	19	18	21	11	0	0	0	0	0	0	0	0	0	0	0	0
Ferritin light chain	P02792	FRIL_HUMAN	1	1	4	3	5	0	1	0	0	0	3	0	2	0	0	0	5	5	5	7	0	0	4	5	5	5	0	0
Fibrinogen beta chain	P02675	FIBB_HUMAN	24	26	1	1	2	24	25	36	41	16	2	23	27	44	44	16	0	0	5	0	0	0	0	0	5	0	0	0
Filaggrin	P20930	FILA_HUMAN	1	1	0	0	0	1	0	1	0	2	0	1	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Filaggrin-2	Q5D862	FILA2_HUMAN	1	1	2	2	0	1	2	1	0	1	0	1	2	1	0	1	0	5	0	0	5	0	0	5	0	0	4	0
Flavin reductase	P30043	BLVRB_HUMAN	0	0	0	0	0	0	0	1	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Fructose-1,6-bisphosphatase 1	P09467	F16P1_HUMAN	0	0	0	0	0	0	2	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Fructose-bisphosphate aldolase B	P05062	ALDOB_HUMAN	25	24	0	0	14	25	44	37	22	5	14	26	40	36	23	3	0	0	0	0	0	0	0	0	0	0	0	0
Galectin-3	P17931	LEG3_HUMAN	4	4	0	0	10	2	4	1	5	2	12	2	3	0	6	3	0	0	0	0	0	0	0	0	0	0	0	0
Galectin-3-binding protein	Q08380	LG3BP_HUMAN	6	6	11	8	12	1	1	8	2	14	11	1	1	9	2	14	0	0	0	7	5	54	0	0	0	0	0	48
Galectin-4	P56470	LEG4_HUMAN	5	5	6	6	6	3	6	6	6	5	6	3	5	6	6	5	10	5	0	14	9	0	8	5	0	12	8	0
Galectin-7	P47929	LEG7_HUMAN	3	3	4	3	0	0	0	0	14	3	0	0	0	0	14	3	0	0	0	21	0	0	0	0	0	17	0	0
Gastric triacylglycerol lipase	P07098	LIPG_HUMAN	15	17	5	5	0	0	3	1	5	82	0	0	3	1	5	94	10	0	0	0	0	22	8	0	0	0	0	21
Gastricsin	P20142	PEPC_HUMAN	5	5	0	0	2	0	2	4	1	21	3	0	3	5	1	20	0	0	0	0	0	0	0	0	0	0	0	0
GDP-mannose 4,6 dehydratase	O60547	GMDS_HUMAN	0	0	0	0	0	0	1	0	1	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Gelsolin	P06396	GELS_HUMAN	2	1	0	0	0	1	1	1	1	5	0	0	0	1	0	3	0	0	0	0	0	0	0	0	0	0	0	0
Glucose-6-phosphate isomerase	P06744	G6PI_HUMAN	7	6	6	5	5	4	7	12	8	5	4	5	5	9	9	5	10	0	0	0	15	11	8	0	0	0	12	9
Glucosidase 2 subunit beta	P14314	GLU2B_HUMAN	1	1	0	0	2	0	0	0	1	0	2	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Glutamyl aminopeptidase	Q07075	AMPE_HUMAN	3	3	2	1	7	0	0	4	4	3	9	0	0	4	3	3	0	0	0	0	9	3	0	0	0	0	4	3
Glutaredoxin-1	P35754	GLRX1_HUMAN	1	1	0	0	0	0	1	1	5	0	0	1	1	1	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Glutathione S-transferase A1	P08263	GSTA1_HUMAN	3	3	0	0	0	9	9	1	0	1	0	8	9	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Glutathione S-transferase omega-1	P78417	GSTO1_HUMAN	2	1	0	0	1	1	2	1	4	1	1	1	2	1	2	1	0	0	0	0	0	0	0	0	0	0	0	0
Glutathione S-transferase P	P09211	GSTP1_HUMAN	6	5	2	2	5	5	6	5	9	6	4	5	4	5	9	5	0	0	0	14	0	0	0	0	0	12	0	0
Glyceraldehyde-3-phosphate dehydrogenase	P04406	G3P_HUMAN	13	12	11	9	7	18	13	4	22	12	6	15	12	4	23	11	10	0	0	21	15	18	8	0	0	17	12	15
Glycerol-3-phosphate dehydrogenase [NAD+], cytoplasmic	P21695	GPDA_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
GTP cyclohydrolase 1 feedback regulatory protein	P30047	GFRP_HUMAN	1	1	0	0	0	1	2	1	1	0	0	1	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Guanine deaminase	Q9Y2T3	GUAD_HUMAN	3	2	0	0	0	5	4	1	4	1	0	3	3	1	3	1	0	0	0	0	0	0	0	0	0	0	0	0
Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-12	Q9UBI6	GBG12_HUMAN	0	0	0	0	0	0	1	0	1	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1	P62873	GBB1_HUMAN	1	1	0	0	0	0	1	4	1	0	0	0	0	4	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Haptoglobin	P00738	HPT_HUMAN	54	50	1	1	9	64	53	44	34	121	9	58	48	41	29	117	5	0	0	0	0	0	4	0	0	0	0	0
Heat shock 70 kDa protein 1A/1B	P08107	HSP71_HUMAN	4	3	2	1	0	3	9	2	6	1	0	3	8	2	6	1	0	0	0	14	0	0	0	0	0	5	0	0
Heat shock cognate 71 kDa protein	P11142	HSP7C_HUMAN	8	7	1	0	0	14	13	7	12	1	0	12	13	7	9	1	0	0	0	7	0	0	0	0	0	0	0	0
Heat shock protein beta-1	P04792	HSPB1_HUMAN	2	2	1	1	0	1	1	1	6	1	0	1	1	1	5	1	0	0	0	7	0	0	0	0	0	5	0	0
Heat shock protein HSP 90-beta	P08238	HS90B_HUMAN	1	1	0	0	0	0	2	0	5	0	0	0	2	0	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Heme-binding protein 1	Q9NRV9	HEBP1_HUMAN	3	3	0	0	0	3	7	4	5	1	0	3	4	4	6	1	0	0	0	0	0	0	0	0	0	0	0	0
Hemoglobin subunit alpha	P69905	HBA_HUMAN	39	36	3	2	10	108	43	27	32	12	7	108	38	24	30	9	0	0	12	0	0	3	0	0	11	0	0	3
Hemoglobin subunit beta	P68871	HBB_HUMAN	64	63	25	22	17	140	61	72	59	32	14	142	60	72	58	30	24	0	77	14	9	25	21	0	58	21	8	24
Hemoglobin subunit delta	P02042	HBD_HUMAN	34	33	21	18	9	78	30	36	33	15	9	78	31	34	33	14	24	0	65	7	9	22	21	0	43	17	8	18
Hemoglobin subunit gamma-2	P69892	HBG2_HUMAN	15	14	9	7	7	31	12	18	13	8	5	28	12	17	13	7	19	0	18	0	5	11	17	0	11	0	4	9

Hemopexin	P02790	HEMO_HUMAN	30	32	2	2	15	40	32	30	31	30	16	45	37	30	33	28	10	0	0	0	0	0	8	0	0	0	0	3
Histidine ammonia-lyase	P42357	HUTH_HUMAN	0	0	0	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Histidine-rich glycoprotein	P04196	HRG_HUMAN	4	4	1	1	0	0	3	1	15	2	0	0	3	1	14	5	0	0	0	0	5	0	0	0	0	0	4	0
Histone H1.3	P16402	H13_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Histone H2A type 1-B/E	P04908	H2A1B_HUMAN	1	1	5	4	0	0	0	0	5	0	0	0	0	0	6	0	0	5	0	14	9	0	0	5	0	12	4	0
Histone H3.1	P68431	H31_HUMAN	1	1	0	0	0	0	3	0	1	0	0	0	2	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Histone H4	P62805	H4_HUMAN	5	4	7	8	0	1	19	0	6	1	0	1	17	0	6	2	0	16	0	14	9	3	0	13	0	17	12	3
HLA class I histocompatibility antigen, A-33 alpha chain	P16190	1A33_HUMAN	1	0	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
HLA class I histocompatibility antigen, B-15 alpha chain	P30464	1B15_HUMAN	1	1	0	0	1	0	0	1	1	2	1	0	0	1	1	2	0	0	0	0	0	0	0	0	0	0	0	0
HLA class II histocompatibility antigen, DR alpha chain	P01903	DRA_HUMAN	1	1	0	0	1	0	0	0	2	1	1	0	0	0	2	1	0	0	0	0	0	0	0	0	0	0	0	0
HLA class II histocompatibility antigen, DRB1-11 beta chain	P20039	2B1B_HUMAN	1	1	0	0	1	1	0	1	1	2	1	1	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Hornerin	Q86YZ3	HORN_HUMAN	2	2	1	1	1	1	3	2	2	1	1	2	2	2	1	1	0	0	0	7	0	0	0	0	0	5	0	0
Hydroxymethylglutaryl-CoA synthase, mitochondrial	P54868	HMCS2_HUMAN	2	2	0	0	0	0	2	0	11	0	0	0	2	0	12	0	0	0	0	0	0	0	0	0	0	0	0	0
Ig alpha-1 chain C region	P01876	IGHA1_HUMAN	191	187	247	234	268	130	132	230	134	254	254	124	124	225	126	267	254	330	120	140	251	387	256	315	110	132	234	354
Ig alpha-2 chain C region	P01877	IGHA2_HUMAN	145	139	290	284	207	94	97	167	111	192	200	89	89	157	107	192	303	381	132	234	357	333	281	369	110	198	427	319
Ig gamma-1 chain C region	P01857	IGHG1_HUMAN	128	121	132	121	64	164	159	133	121	125	62	154	148	123	117	124	168	55	162	183	149	76	153	45	179	144	133	69
Ig gamma-2 chain C region	P01859	IGHG2_HUMAN	67	69	82	68	57	97	75	79	39	55	53	90	81	79	48	60	96	55	132	147	44	18	82	45	132	104	28	15
Ig gamma-3 chain C region	P01860	IGHG3_HUMAN	65	62	78	72	47	81	77	69	52	61	42	78	74	65	50	61	120	33	102	110	77	28	99	35	110	87	72	27
Ig gamma-4 chain C region	P01861	IGHG4_HUMAN	68	70	69	61	47	92	78	82	51	60	43	91	82	80	58	65	96	33	102	124	44	14	87	27	100	87	48	18
Ig heavy chain V-I region HG3	P01743	HV102_HUMAN	4	7	2	1	4	5	1	4	6	5	6	9	3	6	7	10	0	0	0	0	9	0	0	0	0	0	8	0
Ig heavy chain V-II region ARH-77	P06331	HV209_HUMAN	2	3	10	10	2	1	1	1	1	5	5	1	2	1	2	7	0	11	0	14	24	11	0	9	0	12	28	9
Ig heavy chain V-III region BRO	P01766	HV305_HUMAN	26	27	39	33	28	29	19	30	21	29	29	29	19	31	22	31	24	22	5	82	44	57	21	18	11	61	40	48
Ig heavy chain V-III region BUT	P01767	HV306_HUMAN	11	10	7	6	14	11	9	13	8	12	13	10	9	12	7	11	0	0	0	14	15	11	0	0	0	12	12	9
Ig heavy chain V-III region CAM	P01768	HV307_HUMAN	1	1	0	0	1	0	1	0	1	1	0	0	0	1	0	3	0	0	0	0	0	0	0	0	0	0	0	0
Ig heavy chain V-III region GA	P01769	HV308_HUMAN	1	1	3	2	0	1	0	1	0	1	0	1	0	1	0	1	0	0	0	0	15	0	0	0	0	0	12	0
Ig heavy chain V-III region GAL	P01781	HV320_HUMAN	4	4	6	5	7	2	1	6	1	7	6	2	1	6	1	7	5	0	0	7	15	8	4	0	0	5	12	6
Ig heavy chain V-III region TUR	P01779	HV318_HUMAN	11	11	31	25	19	1	7	12	8	16	20	1	8	14	7	16	5	33	18	59	49	22	4	27	21	38	40	18
Ig heavy chain V-III region VH26	P01764	HV303_HUMAN	13	17	25	34	26	5	9	13	8	16	31	5	12	20	12	23	5	16	0	66	29	36	8	35	21	54	57	30
Ig heavy chain V-III region WEA	P01763	HV302_HUMAN	10	10	9	8	12	10	9	11	8	12	12	9	9	12	7	11	0	0	0	14	29	11	0	0	0	12	25	9
Ig kappa chain C region	P01834	IGKC_HUMAN	175	162	342	314	164	155	160	200	116	254	148	143	146	183	112	241	298	369	264	456	395	272	264	333	231	446	358	249
Ig kappa chain V-I region DEE	P01597	KV105_HUMAN	1	1	1	2	1	2	1	0	1	1	2	2	1	0	0	1	0	0	0	0	5	3	0	5	0	0	4	3
Ig kappa chain V-I region HK101 (Fragment)	P01601	KV109_HUMAN	1	2	0	7	1	1	1	0	0	1	3	4	2	1	1	1	0	0	0	0	0	0	0	9	0	17	4	9
Ig kappa chain V-I region HK102 (Fragment)	P01602	KV110_HUMAN	1	3	5	6	2	0	1	1	1	3	5	2	3	2	2	5	5	5	0	7	9	3	4	5	0	12	8	6
Ig kappa chain V-I region Scw	P01609	KV117_HUMAN	2	2	6	6	3	3	2	1	1	2	3	4	2	1	1	2	0	5	0	21	5	3	0	9	0	17	4	6
Ig kappa chain V-II region GM607 (Fragment)	P06309	KV205_HUMAN	7	7	23	19	6	7	6	7	6	9	6	6	5	8	6	9	19	11	5	51	34	18	17	9	5	38	28	15
Ig kappa chain V-III region CLL	P04207	KV308_HUMAN	3	3	4	5	2	3	2	2	2	5	2	3	2	2	2	7	10	0	0	0	0	11	8	5	0	0	4	12
Ig kappa chain V-III region NG9 (Fragment)	P01621	KV303_HUMAN	4	9	9	13	6	3	4	1	2	6	10	8	8	8	6	14	10	11	18	0	5	8	17	9	16	17	12	9
Ig kappa chain V-III region SIE	P01620	KV302_HUMAN	15	15	33	28	14	11	11	19	9	24	13	11	12	19	9	23	34	27	24	59	38	14	29	27	21	45	32	15
Ig kappa chain V-III region VG (Fragment)	P04433	KV309_HUMAN	2	5	5	14	2	1	3	1	2	2	3	4	8	5	6	6	10	5	0	0	5	8	17	13	5	17	16	18
Ig kappa chain V-IV region (Fragment)	P06312	KV401_HUMAN	9	9	28	25	5	10	9	11	6	10	8	9	10	9	6	11	19	27	12	44	38	28	17	27	11	33	36	27
Ig lambda chain V-I region WAH	P04208	LV106_HUMAN	2	3	5	4	2	3	2	2	1	3	3	4	2	5	1	3	5	5	0	7	15	0	4	5	0	5	8	0
Ig lambda chain V-III region LOI	P80748	LV302_HUMAN	5	6	7	8	4	5	6	4	5	8	4	8	4	4	6	9	15	0	0	14	9	3	12	9	0	12	8	6
Ig lambda chain V-III region SH	P01714	LV301_HUMAN	2	2	4	4	1	3	2	1	1	2	1	3	3	1	1	2	5	0	0	0	9	8	4	0	0	0	8	9
Ig lambda-2 chain C regions	P0CG05	LAC2_HUMAN	79	71	151	132	81	95	84	63	48	102	69	87	72	60	45	91	154	165	107	147	231	100	133	144	94	127	206	87
Ig mu heavy chain disease protein	P04220	MUCB_HUMAN	53	52	93	90	82	37	21	59	35	83	79	38	22	57	34	82	105	149	42	0	62	197	95	144	32	5	57	204
IgGFc-binding protein	Q9Y6R7	FCGBP_HUMAN	50	47	162	138	54	23	39	93	35	55	50	22	37	89	35	50	193	116	126	169	188	179	182	90	94	137	165	157
Immunoglobulin J chain	P01591	IGJ_HUMAN	18	18	18	16	18	15	15	27	9	24	18	14	14	25	10	24	24	38	5	0	15	28	17	32	5	0	16	24
Immunoglobulin lambda-like polypeptide 5	B9A064	IGLL5_HUMAN	50	45	94	84	54	60	47	43	28	69	46	53	42	40	23	65	101	100	42	131	111	79	82	90	43	116	105	69
Intelectin-1	Q8WWA0	ITLN1_HUMAN	5	5	18	16	2	11	2	5	5	7	1	9	2	5	5	7	53	5	0	7	20	22	58	0	0	5	12	18
Intelectin-2	Q8WWU7	ITLN2_HUMAN	3	3	10	10	1	2	0	1	6	5	1	2	1	1	6	5	10	0	0	0	38	14	12	0	0	0	36	12
Inter-alpha-trypsin inhibitor heavy chain H1	P19827	ITIH1_HUMAN	2	1	0	0	0	0	3	1	4	1	0	0	2	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Intercellular adhesion molecule 1	P05362	ICAM1_HUMAN	1	1	0	0	0	0	0	0	0	6	0	0	0	0	0	7	0	0	0	0	0	0	0	0	0	0	0	0
Intestinal-type alkaline phosphatase	P09923	PPBI_HUMAN	5	5	79	76	13	1	1	6	6	0	12	1	1	6	6	1	120	160	35	21	97	39	124	153	27	21	97	33
Involucrin	P07476	INVO_HUMAN	1	1	0	0	0	0	0	0	5	0	0	0	0	0	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 1 of Glycerol kinase	P32189-1	GLPK_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 1 of Protein POF1B	Q8WVV4-1	POF1B_HUMAN	1	1	0	0	0	0	0	0	4	1	0	0	0	0	3	1	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 1 of Voltage-dependent anion-selective channel protein 2	P45880-1	VDAC2_HUMAN	1	1	0	0	0	0	0	2	1	0	0	0	0	2	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 10 of Carcinoembryonic antigen-related cell adhesion molecule 1	P13688-10	CEAM1_HUMAN	2	2	5	4	1	0	0	2	1	5	2	0	0	2	1	6	0	11	0	0	5	11	0	9	0	0	4	12
Isoform 10 of Fibronectin	P02751-10	FINC_HUMAN	4	4	0	0	1	8	12	2	2	1	2	6	11	2	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of 14-3-3 protein sigma	P31947-2	1433S_HUMAN	3	2	1	2	0	1	1	0	13	1	0	1	0	0	10	1	0	0	0	7	0	0	0	0	0	12	0	0
Isoform 2 of 3'(2'),5'-bisphosphate nucleotidase 1	O95861-2	BPNT1_HUMAN	1	1	0	0	0	1	1	0	4	0	0	1	1	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Adenylate kinase 2, mitochondrial	P54819-2	KAD2_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Alcohol dehydrogenase 4	P08319-2	ADH4_HUMAN	1	1	0	0	0	2	4	0	0	0	0	2	3	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Annexin A2	P07355-2	ANXA2_HUMAN	11	10	1	0	0	2	6	18	33	7	0	2	5	18	30	6	0	0	0	7	0	0	0	0	0	0	0	0
Isoform 2 of Cadherin-related family member 5	Q9HBB8-2	CDHR5_HUMAN	5	4	0	0	9	1	1	4	7	5	8	1	1	4	7	5	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Catenin alpha-1	P35221-2	CTNA1_HUMAN	1	1	0	0	1	0	0	1	1	0	1	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Clathrin heavy chain 1	Q00610-2	CLH1_HUMAN	2	1	0	0	0	2	4	1	2	0	0	2	3	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Cocaine esterase	O00748-2	EST2_HUMAN	0	0	0	0	0	0	1	0	1	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Creatine kinase U-type, mitochondrial	P12532-2	KCRU_HUMAN	3	3	1	1	1	2	4	4	7	2	1	2	3	2	7	2	0	0	0	0	5	0	0	0	0	0	4	0
Isoform 2 of Cytosol aminopeptidase	P28838-2	AMPL_HUMAN	5	5	8	8	8	3	4	8	6	1	6	3	4	9	5	1	10	5	5	7	15	3	12	9	5	5	12	3
Isoform 2 of Delta-aminolevulinic acid dehydratase	P13716-2	HEM2_HUMAN	1	0	0	0	0	1	1	0	1	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Elongation factor 1-delta	P29692-2	EF1D_HUMAN	1	1	0	0	1	1	2	1	1	1	1	1	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Ester hydrolase C11orf54	Q9H0W9-2	CK054_HUMAN	2	2	2	2	2	0	2	1	5	1	2	0	2	1	3	1	0	0	0	0	9	0	0	0	0	0	12	0
Isoform 2 of Fatty aldehyde dehydrogenase	P51648-2	AL3A2_HUMAN	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Fibrinogen alpha chain	P02671-2	FIBA_HUMAN	7	7	0	0	0	3	9	14	8	7	0	5	9	14	8	7	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Filamin-B	O75369-2	FLNB_HUMAN	5	6	0	0	12	3	6	8	2	1	12	3	7	9	3	2	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Gastric intrinsic factor	P27352-2	IF_HUMAN	3	2	7	6	6	0	0	0	1	8	6	0	0	0	0	8	5	22	0	0	5	11	4	18	0	0	4	9

Isoform 2 of Glyoxalase domain-containing protein 4	Q9HC38-2	GLOD4_HUMAN	1	0	0	0	0	1	2	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of G-protein coupled receptor 126	Q86SQ4-2	GP126_HUMAN	1	1	0	0	0	0	0	1	0	6	0	0	0	1	0	3	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Heat shock protein HSP 90-alpha	P07900-2	HS90A_HUMAN	2	2	0	0	0	1	4	0	5	0	0	1	7	0	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial	Q16836-2	HCDH_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Ig mu chain C region	P01871-2	IGHM_HUMAN	80	77	162	153	133	53	29	91	53	121	127	52	27	89	51	117	164	220	60	44	126	357	148	211	48	54	112	346
Isoform 2 of Inorganic pyrophosphatase 2, mitochondrial	Q9H2U2-2	IPYR2_HUMAN	1	1	0	0	0	1	1	1	2	0	0	1	1	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4	Q14624-2	ITIH4_HUMAN	3	3	0	0	0	7	4	5	2	2	0	5	3	4	2	2	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Lactoylglutathione lyase	Q04760-2	LGUL_HUMAN	1	1	0	0	1	1	0	1	1	1	0	2	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Lambda-crystallin homolog	Q9Y2S2-2	CRYL1_HUMAN	0	0	0	0	0	0	2	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Long-chain-fatty-acid--CoA ligase 5	Q9ULC5-3	ACSL5_HUMAN	5	5	0	0	0	10	10	5	6	0	0	9	10	4	6	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Medium-chain specific acyl-CoA dehydrogenase, mitochondrial	P11310-2	ACADM_HUMAN	6	5	5	4	12	3	2	5	13	0	9	3	3	4	9	0	5	0	0	7	15	0	4	0	0	5	12	0
Isoform 2 of NADH dehydrogenase [ubiquinone] flavoprotein 1, mitochondrial	P49821-2	NDUV1_HUMAN	0	0	0	0	0	0	0	1	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Neutral ceramidase	Q9NR71-2	ASAH2_HUMAN	4	4	0	0	15	0	1	6	1	1	13	0	1	6	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Nicotinate phosphoribosyltransferase	Q6XQN6-2	PNCB_HUMAN	0	0	0	0	0	0	0	0	2	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Nucleoside diphosphate kinase A	P15531-2	NDKA_HUMAN	1	1	0	0	1	1	1	1	2	0	1	1	1	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Proteasome subunit beta type-8	P28062-2	PSB8_HUMAN	1	1	0	0	1	1	2	1	1	1	0	1	2	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Protein disulfide-isomerase A6	Q15084-2	PDIA6_HUMAN	1	1	0	0	1	0	0	2	2	0	2	0	0	2	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Protein JTB	O76095-2	JTB_HUMAN	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Selenium-binding protein 1	Q13228-2	SBP1_HUMAN	2	2	2	2	2	2	2	4	2	1	2	2	2	5	2	1	0	0	0	14	0	0	0	0	0	12	0	0
Isoform 2 of Serine hydroxymethyltransferase, cytosolic	P34896-2	GLYC_HUMAN	2	2	0	0	1	0	4	1	5	0	1	0	4	1	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Sodium/potassium-transporting ATPase subunit beta-1	P05026-2	AT1B1_HUMAN	2	1	0	0	1	0	0	4	4	0	1	0	0	5	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Tripeptidyl-peptidase 1	O14773-2	TPP1_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Tyrosine-protein phosphatase non-receptor type substrate 1	P78324-2	SHPS1_HUMAN	1	1	0	0	0	0	0	0	0	3	0	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Valacyclovir hydrolase	Q86WA6-2	BPHL_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Very long-chain specific acyl-CoA dehydrogenase, mitochondrial	P49748-2	ACADV_HUMAN	12	11	0	0	18	5	12	12	24	2	16	5	12	12	19	2	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Vesicle-associated membrane protein-associated protein A	Q9P0L0-2	VAPA_HUMAN	1	1	0	0	0	1	1	0	2	0	0	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 2 of Xaa-Pro aminopeptidase 1	Q9NQW7-2	XPP1_HUMAN	1	1	0	0	4	2	1	0	1	0	3	2	2	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 3 of 4F2 cell-surface antigen heavy chain	P08195-3	4F2_HUMAN	1	1	0	0	3	0	0	1	1	0	3	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 3 of Apoptosis-inducing factor 1, mitochondrial	O95831-3	AIFM1_HUMAN	1	1	0	0	0	0	2	1	1	1	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 3 of Deleted in malignant brain tumors 1 protein	Q9UGM3-3	DMBT1_HUMAN	29	29	102	103	40	21	17	25	29	44	41	22	16	21	27	47	71	67	42	89	73	268	70	59	37	54	69	328
Isoform 3 of Heterogeneous nuclear ribonucleoprotein K	P61978-3	HNRPK_HUMAN	1	1	0	0	0	0	1	0	2	0	0	0	1	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 8 of Plectin	Q15149-8	PLEC_HUMAN	1	1	0	0	0	1	1	0	1	0	0	0	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Isoform 8 of Titin	Q8WZ42-8	TITIN_HUMAN	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Isoform Alpha of Pancreatic secretory granule membrane major glycoprotein GP2	P55259-3	GP2_HUMAN	82	78	272	264	200	112	94	36	49	2	191	106	92	32	46	3	413	535	221	162	222	76	421	531	215	127	225	63
Isoform B of Trypsin-3	P35030-2	TRY3_HUMAN	33	39	28	117	48	30	32	38	39	8	48	38	40	46	48	15	34	38	12	37	24	22	54	104	53	335	93	60
Isoform Beta of Caspase-6	P55212-2	CASP6_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform C of Ketohexokinase	P50053-2	KHK_HUMAN	1	1	0	0	0	1	2	0	2	0	0	1	2	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform Cytoplasmic of Fumarate hydratase, mitochondrial	P07954-2	FUMH_HUMAN	0	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform Cytoplasmic+peroxisomal of Peroxiredoxin-5, mitochondrial	P30044-2	PRDX5_HUMAN	0	0	0	0	0	0	2	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform Gamma-A of Fibrinogen gamma chain	P02679-2	FIBG_HUMAN	22	20	0	0	0	19	24	39	32	16	0	19	19	38	29	17	0	0	0	0	0	0	0	0	0	0	0	0
Isoform H7 of Myeloperoxidase	P05164-3	PERM_HUMAN	4	4	9	8	0	0	0	0	0	23	0	0	0	0	0	23	10	0	0	0	0	43	12	0	0	0	0	37
Isoform LMP2.S of Proteasome subunit beta type-9	P28065-2	PSB9_HUMAN	1	1	0	0	1	1	2	1	1	1	1	1	2	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Isoform LMW of Kininogen-1	P01042-2	KNG1_HUMAN	3	2	0	0	0	1	4	0	4	7	0	1	3	0	2	7	0	0	0	0	0	0	0	0	0	0	0	0
Isoform PSMA-7 of Glutamate carboxypeptidase 2	Q04609-7	FOLH1_HUMAN	1	1	0	0	1	0	0	1	1	3	1	0	0	0	1	3	0	0	0	0	0	0	0	0	0	0	0	0
Isoform Sap-mu-6 of Proactivator polypeptide	P07602-2	SAP_HUMAN	1	1	0	0	0	0	0	0	0	3	0	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0
Isoform Smooth muscle of Myosin light polypeptide 6	P60660-2	MYL6_HUMAN	2	2	0	0	0	2	4	1	6	0	0	3	4	0	6	0	0	0	0	0	0	0	0	0	0	0	0	0
Isoform Somatic-2 of Angiotensin-converting enzyme	P12821-2	ACE_HUMAN	3	4	0	0	5	0	0	1	7	7	6	0	0	0	7	8	0	0	0	0	0	0	0	0	0	0	0	0
Junction plakoglobin	P14923	PLAK_HUMAN	2	2	2	2	0	1	2	1	8	2	0	1	2	0	8	2	0	0	0	0	0	11	0	0	0	0	0	9
Junctional adhesion molecule A	Q9Y624	JAM1_HUMAN	3	4	1	1	2	0	0	4	4	5	3	0	0	5	6	7	5	0	0	0	0	0	4	0	0	0	0	0
Kallikrein-1	P06870	KLK1_HUMAN	3	3	0	0	8	1	2	1	2	6	7	1	2	1	2	6	0	0	0	0	0	0	0	0	0	0	0	0
Kallistatin	P29622	KAIN_HUMAN	1	1	0	0	2	2	1	1	2	0	2	2	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Lactase-phlorizin hydrolase	P09848	LPH_HUMAN	21	18	6	5	43	10	6	35	26	5	40	8	4	31	24	2	0	0	5	14	15	3	0	0	5	12	12	3
Lactoperoxidase	P22079	PERL_HUMAN	2	2	2	2	0	0	0	0	0	10	0	0	0	0	0	10	0	0	0	0	0	11	0	0	0	0	0	9
Lactotransferrin	P02788	TRFL_HUMAN	39	35	20	17	6	0	0	1	0	227	5	0	0	1	0	201	5	0	0	0	0	114	4	0	0	0	0	97
Lamin-B1	P20700	LMNB1_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Leucine-rich alpha-2-glycoprotein	P02750	A2GL_HUMAN	3	3	0	0	2	5	4	4	1	2	2	3	4	4	1	2	0	0	0	0	0	0	0	0	0	0	0	0
Leukocyte elastase inhibitor	P30740	ILEU_HUMAN	9	8	0	0	17	3	10	8	8	10	15	3	9	7	7	9	0	0	0	0	0	0	0	0	0	0	0	0
Leukotriene A-4 hydrolase	P09960	LKHA4_HUMAN	1	0	0	0	0	2	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
LIM and SH3 domain protein 1	Q14847	LASP1_HUMAN	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Lipocalin-1	P31025	LCN1_HUMAN	3	4	0	0	0	0	0	0	0	16	0	0	0	0	0	23	0	0	0	0	0	0	0	0	0	0	0	0
Lithostathine-1-alpha	P05451	REG1A_HUMAN	22	22	35	32	22	19	15	29	24	21	24	18	16	31	27	18	53	27	30	73	15	11	54	22	32	61	12	9
L-lactate dehydrogenase A chain	P00338	LDHA_HUMAN	5	5	1	1	0	0	23	1	5	2	0	0	22	1	3	2	0	0	0	7	0	0	0	0	0	5	0	0
Long palate, lung and nasal epithelium carcinoma-associated protein 1	Q8TDL5	LPLC1_HUMAN	4	5	3	4	0	0	0	0	0	22	0	0	0	0	0	27	5	0	0	0	0	14	4	0	0	0	0	18
Lysozyme C	P61626	LYSC_HUMAN	22	27	2	2	9	8	23	25	29	36	12	9	28	28	36	49	0	0	0	0	0	14	0	0	0	0	0	12
Macrophage migration inhibitory factor	P14174	MIF_HUMAN	2	1	0	0	0	2	2	1	2	2	0	2	1	1	1	2	0	0	0	0	0	0	0	0	0	0	0	0
Malate dehydrogenase, cytoplasmic	P40925	MDHC_HUMAN	5	4	0	0	0	5	9	4	5	5	1	4	9	4	3	3	0	0	0	0	0	0	0	0	0	0	0	0
Malate dehydrogenase, mitochondrial	P40926	MDHM_HUMAN	22	19	0	0	14	23	32	30	25	7	12	19	26	28	23	7	0	0	0	0	0	0	0	0	0	0	0	0
Maltase-glucoamylase, intestinal	O43451	MGA_HUMAN	9	8	12	11	24	2	6	5	14	2	22	1	4	4	12	3	5	33	0	0	24	8	4	32	0	0	21	6
Membrane-associated progesterone receptor component 2	O15173	PGRC2_HUMAN	1	1	0	0	0	1	0	2	1	0	0	1	0	2	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Meprin A subunit alpha	Q16819	MEP1A_HUMAN	5	6	8	5	9	0	1	6	6	7	12	0	1	8	5	7	0	5	0	7	15	18	0	5	0	0	12	15
Meprin A subunit beta	Q16820	MEP1B_HUMAN	3	2	0	0	10	1	1	4	2	1	6	1	0	4	2	1	0	0	0	0	0	0	0	0	0	0	0	0
Microsomal glutathione S-transferase 3	O14880	MGST3_HUMAN	1	1	1	1	0	0	0	1	5	1	0	0	0	1	5	1	0	0	0	0	5	0	0	0	0	0	4	0
Microsomal triglyceride transfer protein large subunit	P55157	MTP_HUMAN	6	6	0	0	0	8	25	0	5	0	0	8	24	0	6	0	0	0	0	0	0	0	0	0	0	0	0	0
Mucin-13	Q9H3R2	MUC13_HUMAN	2	3	0	0	1	1	1	6	2	3	0	1	1	6	3	5	0	0	0	0	0	0	0	0	0	0	0	0
Mucin-2	Q02817	MUC2_HUMAN	7	7	8	7	2	2	2	22	5	8	3	2	2	19	5	8	10	11	5	0	9	11	8	9	5	0	8	9
Mucin-5AC (Fragments)	P98088	MUC5A_HUMAN	42	40	4	5	24	0	14	6	11	195	25	0	13	6	10	185	0	0	0	0	0	22	0	0	0	0	0	30
Mucin-5B	Q9HC84	MUC5B_HUMAN	25	23	31	31	2	0	1	1	1	144	3	0	1	1	1	134	24	0	0	0	0	164	25	0	0	0	0	162

Mucin-6	Q6W4X9	MUC6_HUMAN	29	27	10	16	29	5	17	54	31	36	29	3	16	51	27	38	10	5	0	7	24	14	12	18	0	5	28	33
Multimerin-1	Q13201	MMRN1_HUMAN	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Myeloblastin	P24158	PRTN3_HUMAN	1	1	1	1	0	0	0	0	0	5	0	0	0	0	0	5	0	0	0	0	0	3	0	0	0	0	0	3
Myosin regulatory light chain 12B	O14950	ML12B_HUMAN	2	2	0	0	1	1	2	2	5	0	0	0	2	2	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Myosin-9	P35579	MYH9_HUMAN	1	1	1	0	0	0	1	0	2	2	0	0	0	0	2	2	0	0	0	7	0	0	0	0	0	0	0	0
Na(+)/H(+) exchange regulatory cofactor NHE-RF1	O14745	NHRF1_HUMAN	1	1	0	0	0	1	3	1	2	0	0	1	3	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
NADH dehydrogenase [ubiquinone] iron-sulfur protein 3, mitochondrial	O75489	NDUS3_HUMAN	0	0	0	0	1	0	0	1	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
NADPH--cytochrome P450 reductase	P16435	NCPR_HUMAN	1	0	0	0	0	0	2	0	1	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Neprilysin	P08473	NEP_HUMAN	8	7	17	15	21	0	0	10	7	7	18	0	0	11	6	7	0	38	0	0	9	57	0	32	0	0	8	48
Neuroblast differentiation-associated protein AHNAK	Q09666	AHNK_HUMAN	1	0	0	0	0	0	0	0	4	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Neutrophil defensin 1	P59665	DEF1_HUMAN	7	6	0	0	4	0	1	6	1	28	3	0	1	6	1	24	0	0	0	0	0	0	0	0	0	0	0	0
Neutrophil elastase	P08246	ELNE_HUMAN	1	1	1	1	0	0	0	0	0	5	0	0	0	0	0	5	0	0	0	0	0	3	0	0	0	0	0	3
Neutrophil gelatinase-associated lipocalin	P80188	NGAL_HUMAN	4	3	6	6	1	0	0	1	1	18	1	0	0	1	1	17	19	0	0	0	5	14	17	0	0	0	4	12
Olfactomedin-4	Q6UX06	OLFM4_HUMAN	25	25	99	98	40	7	4	39	24	38	39	6	3	38	23	39	58	94	0	110	145	189	66	85	0	111	141	187
Ornithine aminotransferase, mitochondrial	P04181	OAT_HUMAN	4	4	0	0	3	3	14	0	5	1	3	3	14	0	5	1	0	0	0	0	0	0	0	0	0	0	0	0
Pancreatic alpha-amylase	P04746	AMYP_HUMAN	189	198	438	464	220	203	177	136	141	254	226	213	189	148	152	259	365	243	700	557	348	415	392	256	688	594	386	469
Pancreatic lipase-related protein 1	P54315	LIPR1_HUMAN	9	8	4	8	15	12	10	8	6	0	17	10	9	7	6	0	5	5	5	0	9	0	12	18	11	0	8	0
Pancreatic lipase-related protein 2	P54317	LIPR2_HUMAN	32	32	32	32	75	47	9	25	33	2	79	47	9	25	30	2	34	67	42	14	29	8	37	67	48	5	25	12
Pancreatic triacylglycerol lipase	P16233	LIPP_HUMAN	247	273	189	206	425	294	234	230	244	53	484	325	260	251	266	53	216	325	311	59	159	61	235	360	331	54	185	72
Pantetheinase	O95497	VNN1_HUMAN	4	4	15	11	4	0	0	4	7	7	3	0	0	4	7	7	0	44	0	7	38	0	0	35	0	5	28	0
Pepsin A	P00790	PEPA_HUMAN	0	1	0	0	0	0	0	0	0	1	2	2	1	1	0	2	0	0	0	0	0	0	0	0	0	0	0	0
Peptidyl-prolyl cis-trans isomerase B	P23284	PPIB_HUMAN	1	1	0	0	0	1	4	0	1	1	0	1	3	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Periplakin	O60437	PEPL_HUMAN	0	0	0	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Peroxiredoxin-1	Q06830	PRDX1_HUMAN	7	7	0	0	1	12	11	6	9	3	1	11	12	7	10	3	0	0	0	0	0	0	0	0	0	0	0	0
Peroxiredoxin-2	P32119	PRDX2_HUMAN	4	3	0	0	0	3	12	1	6	1	0	2	11	1	6	0	0	0	0	0	0	0	0	0	0	0	0	0
Peroxiredoxin-4	Q13162	PRDX4_HUMAN	3	2	0	0	2	3	4	4	2	1	2	3	3	4	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Peroxisomal acyl-coenzyme A oxidase 1	Q15067	ACOX1_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Peroxisomal multifunctional enzyme type 2	P51659	DHB4_HUMAN	3	3	0	0	2	2	3	4	5	1	1	2	3	4	5	1	0	0	0	0	0	0	0	0	0	0	0	0
Phenazine biosynthesis-like domain-containing protein	P30039	PBLD_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Phosphatidylethanolamine-binding protein 1	P30086	PEBP1_HUMAN	12	10	0	0	4	14	18	13	13	7	4	12	15	12	13	6	0	0	0	0	0	0	0	0	0	0	0	0
Phosphoenolpyruvate carboxykinase [GTP], mitochondrial	Q16822	PCKGM_HUMAN	6	6	0	0	0	1	32	0	1	0	0	1	31	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Phosphoglycerate kinase 1	P00558	PGK1_HUMAN	4	3	0	0	1	5	11	0	4	0	1	5	10	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0
Phosphoglycerate mutase 1	P18669	PGAM1_HUMAN	2	3	0	0	1	2	3	1	6	0	1	3	4	1	6	0	0	0	0	0	0	0	0	0	0	0	0	0
Phospholipase A2	P04054	PA21B_HUMAN	17	20	23	31	30	24	13	11	19	2	39	31	16	12	19	2	19	16	60	30	15	0	25	18	84	33	28	0
Phospholipase A2, membrane associated	P14555	PA2GA_HUMAN	2	2	0	0	4	1	0	1	1	5	3	1	0	0	1	6	0	0	0	0	0	0	0	0	0	0	0	0
Phospholysine phosphohistidine inorganic pyrophosphate phosphatase	Q9H008	LHPP_HUMAN	0	0	0	0	0	1	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Phosphomannomutase 2	O15305	PMM2_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Plasma protease C1 inhibitor	P05155	IC1_HUMAN	11	12	0	0	6	13	10	13	9	12	5	14	12	14	12	15	0	0	0	0	0	0	0	0	0	0	0	0
Plasma serine protease inhibitor	P05154	IPSP_HUMAN	2	2	0	0	1	1	1	2	4	1	1	1	1	4	2	1	0	0	0	0	0	0	0	0	0	0	0	0
Plasminogen	P00747	PLMN_HUMAN	2	2	0	0	2	4	2	1	1	0	2	3	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Plastin-1	Q14651	PLSI_HUMAN	4	4	0	0	2	3	7	1	7	1	2	3	7	1	7	1	0	0	0	0	0	0	0	0	0	0	0	0
Polymeric immunoglobulin receptor	P01833	PIGR_HUMAN	160	159	104	89	196	72	82	184	109	315	185	70	74	187	110	326	81	138	18	51	87	251	75	104	11	33	80	229
Prelamin-A/C	P02545	LMNA_HUMAN	2	1	0	0	0	1	2	0	4	2	0	1	2	0	2	1	0	0	0	0	0	0	0	0	0	0	0	0
Profilin-1	P07737	PROF1_HUMAN	2	2	0	0	0	0	4	0	5	0	0	0	4	0	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Prolactin-inducible protein	P12273	PIP_HUMAN	7	7	2	3	2	0	0	0	0	42	2	0	0	0	0	40	0	0	0	0	0	14	0	0	0	0	0	18
Prostaglandin reductase 1	Q14914	PTGR1_HUMAN	1	0	0	0	1	0	0	0	5	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Prostaglandin-H2 D-isomerase	P41222	PTGDS_HUMAN	1	1	0	0	5	0	0	0	0	0	4	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Prostasin	Q16651	PRSS8_HUMAN	2	4	0	0	3	1	0	1	4	3	5	1	0	4	5	6	0	0	0	0	0	0	0	0	0	0	0	0
Prostate stem cell antigen	O43653	PSCA_HUMAN	1	1	0	0	2	0	0	0	0	2	3	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0
Proteasome subunit alpha type-2	P25787	PSA2_HUMAN	1	1	0	0	0	0	1	1	1	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Proteasome subunit alpha type-4	P25789	PSA4_HUMAN	0	0	0	0	0	1	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Proteasome subunit alpha type-6	P60900	PSA6_HUMAN	1	1	0	0	1	2	2	1	1	0	1	2	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Proteasome subunit beta type-1	P20618	PSB1_HUMAN	1	1	0	0	0	1	1	1	1	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Proteasome subunit beta type-2	P49721	PSB2_HUMAN	1	1	2	2	0	1	1	1	1	0	0	1	1	1	1	0	5	0	0	0	5	3	4	0	0	0	4	3
Proteasome subunit beta type-3	P49720	PSB3_HUMAN	1	1	0	0	0	1	0	2	1	0	0	1	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Protein AMBP	P02760	AMBP_HUMAN	4	3	0	0	1	3	2	1	5	13	1	2	2	1	3	11	0	0	0	0	0	0	0	0	0	0	0	0
Protein disulfide-isomerase	P07237	PDIA1_HUMAN	9	9	1	1	0	14	17	8	16	1	1	11	17	6	15	1	0	0	0	7	0	0	0	0	0	5	0	0
Protein disulfide-isomerase A3	P30101	PDIA3_HUMAN	2	1	0	0	0	3	4	0	2	0	0	2	4	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Protein FAM3C	Q92520	FAM3C_HUMAN	2	2	0	0	2	1	2	2	2	0	2	1	2	4	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Protein Plunc	Q9NP55	PLUNC_HUMAN	5	5	7	6	0	0	0	0	0	29	0	0	0	0	0	32	0	0	0	0	0	39	0	0	0	0	0	33
Protein S100-A11	P31949	S10AB_HUMAN	1	1	0	0	0	0	0	0	2	6	0	0	0	0	2	6	0	0	0	0	0	0	0	0	0	0	0	0
Protein S100-A12	P80511	S10AC_HUMAN	0	0	0	0	0	0	0	0	0	2	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Protein S100-A6	P06703	S10A6_HUMAN	2	2	0	0	2	0	2	4	4	2	2	0	2	4	2	2	0	0	0	0	0	0	0	0	0	0	0	0
Protein S100-A7	P31151	S10A7_HUMAN	3	2	9	7	1	3	2	1	1	8	1	1	2	1	1	8	10	5	12	14	5	8	8	5	5	12	4	6
Protein S100-A8	P05109	S10A8_HUMAN	6	6	6	6	1	2	2	1	1	30	1	1	2	0	1	28	0	0	0	7	0	28	0	0	0	5	0	30
Protein S100-A9	P06702	S10A9_HUMAN	8	7	18	17	2	1	0	1	5	36	3	1	0	1	3	34	10	11	0	30	5	50	8	13	0	21	4	54
Protein S100-G	P29377	S100G_HUMAN	1	1	0	0	0	2	4	1	0	1	0	3	3	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Protein S100-P	P25815	S100P_HUMAN	1	1	0	0	0	0	0	0	0	5	0	0	0	0	0	5	0	0	0	0	0	0	0	0	0	0	0	0
Protein TFG	Q92734	TFG_HUMAN	0	0	0	0	0	0	2	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Protein-glutamine gamma-glutamyltransferase E	Q08188	TGM3_HUMAN	3	3	0	0	0	0	0	1	0	17	0	0	0	1	0	17	0	0	0	0	0	0	0	0	0	0	0	0
Purine nucleoside phosphorylase	P00491	PNPH_HUMAN	1	1	0	0	0	0	4	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Puromycin-sensitive aminopeptidase	P55786	PSA_HUMAN	1	1	0	0	0	1	3	0	1	0	0	1	3	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Putative elongation factor 1-alpha-like 3	Q5VTE0	EF1A3_HUMAN	1	1	0	0	0	0	0	0	4	0	0	0	0	0	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Putative gamma-glutamyltranspeptidase 3	A6NGU5	GGT3_HUMAN	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0
Quinone oxidoreductase	Q08257	QOR_HUMAN	1	1	0	0	0	2	2	1	1	0	0	2	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Rab GDP dissociation inhibitor beta	P50395	GDIB_HUMAN	3	2	0	0	0	4	4	5	1	1	0	4	3	5	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Ras GTPase-activating-like protein IQGAP1	P46940	IQGA1_HUMAN	0	0	0	0	0	0	0	0	1	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0

Ras GTPase-activating-like protein IQGAP2	Q13576	IQGA2_HUMAN	1	1	0	0	0	0	1	1	1	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Ras-related protein Rab-11A	P62491	RB11A_HUMAN	1	1	0	0	0	2	2	1	1	0	0	1	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Ras-related protein Rab-1A	P62820	RAB1A_HUMAN	3	3	0	0	0	3	7	1	5	0	0	2	7	1	6	0	0	0	0	0	0	0	0	0	0	0	0	0
Ras-related protein Rab-2A	P61019	RAB2A_HUMAN	0	0	0	0	0	0	0	0	1	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Ras-related protein Rab-5C	P51148	RAB5C_HUMAN	1	1	0	0	0	0	1	1	1	0	0	0	2	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Ras-related protein Rab-8B	Q92930	RAB8B_HUMAN	1	1	0	0	0	1	4	1	2	0	0	1	3	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0
Regenerating islet-derived protein 3-alpha	Q06141	REG3A_HUMAN	16	19	2	2	13	13	8	13	32	16	17	15	7	12	43	18	5	0	0	0	5	3	4	0	0	0	4	3
Retinal dehydrogenase 1	P00352	AL1A1_HUMAN	5	5	0	0	1	4	15	4	8	0	1	3	14	4	8	0	0	0	0	0	0	0	0	0	0	0	0	0
Retinol-binding protein 2	P50120	RET2_HUMAN	2	2	0	0	0	2	6	1	2	1	0	2	5	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Retinol-binding protein 4	P02753	RET4_HUMAN	3	3	0	0	1	2	2	1	6	5	0	1	2	1	6	5	0	0	0	0	0	0	0	0	0	0	0	0
Rho GDP-dissociation inhibitor 1	P52565	GDIR1_HUMAN	1	1	0	0	1	1	2	1	2	0	1	1	2	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Rho GDP-dissociation inhibitor 2	P52566	GDIR2_HUMAN	1	0	0	0	0	1	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0
Ribonuclease pancreatic	P07998	RNAS1_HUMAN	1	1	0	0	0	1	2	1	0	0	0	1	3	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Ribonuclease T2	O00584	RNT2_HUMAN	1	0	0	0	1	0	0	0	0	2	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Ribonuclease UK114	P52758	UK114_HUMAN	2	1	0	0	1	1	0	4	2	1	1	1	0	2	2	1	0	0	0	0	0	0	0	0	0	0	0	0
Secretoglobin family 3A member 1	Q96QR1	SG3A1_HUMAN	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Secretoglobin family 3A member 2	Q96PL1	SG3A2_HUMAN	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Serpin B3	P29508	SPB3_HUMAN	8	8	9	8	0	0	0	0	39	9	1	1	1	0	35	10	0	0	0	51	0	0	0	0	0	45	0	0
Serpin B4	P48594	SPB4_HUMAN	6	7	9	6	0	0	0	0	31	7	1	1	1	0	31	7	0	0	0	51	0	0	0	0	0	33	0	0
Serpin B6	P35237	SPB6_HUMAN	6	5	7	5	11	5	4	5	8	2	9	4	4	4	7	2	0	5	5	7	24	0	0	5	5	5	16	0
Serum albumin	P02768	ALBU_HUMAN	943	946	327	340	489	1283	1275	1007	980	621	465	1291	1298	1018	978	624	399	193	538	302	188	340	455	189	594	259	217	324
Serum amyloid P-component	P02743	SAMP_HUMAN	2	2	0	0	1	3	6	1	0	0	1	3	7	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Short palate, lung and nasal epithelium carcinoma-associated protein 2	Q96DR5	SPLC2_HUMAN	2	2	0	0	0	0	0	0	0	9	0	0	0	0	0	13	0	0	0	0	0	0	0	0	0	0	0	0
Short-chain specific acyl-CoA dehydrogenase, mitochondrial	P16219	ACADS_HUMAN	2	2	0	0	2	1	1	4	4	0	3	1	2	4	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Sodium/potassium-transporting ATPase subunit alpha-1	P05023	AT1A1_HUMAN	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sorbitol dehydrogenase	Q00796	DHSO_HUMAN	0	1	0	0	0	0	2	0	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Spectrin beta chain, brain 1	Q01082	SPTB2_HUMAN	1	1	0	0	0	1	1	1	2	0	0	0	1	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Stress-70 protein, mitochondrial	P38646	GRP75_HUMAN	1	0	0	0	0	0	3	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial	P31040	DHSA_HUMAN	1	1	0	0	2	0	1	0	2	0	2	0	2	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Succinyl-CoA ligase [GDP-forming] subunit alpha, mitochondrial	P53597	SUCA_HUMAN	1	1	0	0	0	1	1	0	1	0	0	1	2	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sucrase-isomaltase, intestinal	P14410	SUIS_HUMAN	48	44	98	80	65	25	22	75	58	42	61	21	17	67	57	40	58	116	114	147	87	68	54	94	105	99	69	57
Sulfhydryl oxidase 1	O00391	QSOX1_HUMAN	0	1	0	0	0	0	0	0	0	2	0	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0
Superoxide dismutase [Cu-Zn]	P00441	SODC_HUMAN	5	5	2	1	4	5	4	7	4	5	4	4	4	7	5	3	0	0	0	14	0	0	0	0	0	5	0	0
Superoxide dismutase [Mn], mitochondrial	P04179	SODM_HUMAN	2	1	0	0	2	1	1	1	2	2	1	1	1	1	2	2	0	0	0	0	0	0	0	0	0	0	0	0
SWI/SNF complex subunit SMARCC1	Q92922	SMRC1_HUMAN	0	1	0	0	0	0	1	0	0	0	1	1	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Syncollin	Q0VAF6	SYCN_HUMAN	1	2	4	5	1	3	2	1	1	0	1	4	3	1	2	0	10	0	12	0	0	0	12	0	16	0	0	0
Tetranectin	P05452	TETN_HUMAN	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Tetraspanin-8	P19075	TSN8_HUMAN	1	1	0	0	1	0	1	2	1	2	1	0	0	1	1	2	0	0	0	0	0	0	0	0	0	0	0	0
Tetratricopeptide repeat protein 38	Q5R3I4	TTC38_HUMAN	1	1	0	0	1	1	2	1	1	0	1	1	1	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Thioredoxin domain-containing protein 17	Q9BRA2	TXD17_HUMAN	1	1	0	0	0	0	2	0	2	0	0	0	2	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Thioredoxin domain-containing protein 5	Q8NBS9	TXND5_HUMAN	0	0	0	0	0	1	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Thioredoxin-dependent peroxide reductase, mitochondrial	P30048	PRDX3_HUMAN	3	3	0	0	0	2	7	2	4	1	0	2	7	2	3	2	0	0	0	0	0	0	0	0	0	0	0	0
Thymidine phosphorylase	P19971	TYPH_HUMAN	1	0	0	0	0	0	1	0	4	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Thyroxine-binding globulin	P05543	THBG_HUMAN	2	1	0	0	2	3	1	0	2	1	2	2	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
Transaldolase	P37837	TALDO_HUMAN	1	1	0	0	3	1	1	0	1	1	3	1	2	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Transgelin-2	P37802	TAGL2_HUMAN	2	2	0	0	2	2	3	0	5	1	2	3	3	0	3	1	0	0	0	0	0	0	0	0	0	0	0	0
Transitional endoplasmic reticulum ATPase	P55072	TERA_HUMAN	1	1	0	0	0	1	1	0	6	0	0	1	1	0	6	0	0	0	0	0	0	0	0	0	0	0	0	0
Transketolase	P29401	TKT_HUMAN	4	4	3	2	7	2	1	6	2	5	6	2	1	6	2	5	0	0	0	0	9	8	0	0	0	0	8	6
Transthyretin	P02766	TTHY_HUMAN	8	10	13	11	6	12	13	2	8	8	6	14	16	4	9	10	19	5	24	14	15	3	17	5	21	5	12	3
Trefoil factor 1	P04155	TFF1_HUMAN	1	1	0	5	1	0	0	0	0	2	1	0	0	0	0	2	0	0	0	0	0	0	0	9	0	5	0	18
Trefoil factor 2	Q03403	TFF2_HUMAN	6	6	31	26	6	3	7	8	2	12	5	3	7	8	2	11	24	33	24	66	29	11	17	27	27	50	25	9
Trefoil factor 3	Q07654	TFF3_HUMAN	0	1	0	0	0	0	0	0	0	0	1	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0
Tricarboxylate transport protein, mitochondrial	P53007	TXTP_HUMAN	1	1	0	0	0	0	0	2	1	0	0	0	0	2	1	0	0	0	0	0	0	0	0	0	0	0	0	0
Trifunctional enzyme subunit alpha, mitochondrial	P40939	ECHA_HUMAN	0	0	0	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Trifunctional enzyme subunit beta, mitochondrial	P55084	ECHB_HUMAN	0	0	0	0	0	0	0	0	2	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
Triosephosphate isomerase	P60174	TPIS_HUMAN	28	26	3	3	37	29	41	25	21	15	36	25	39	22	20	15	5	0	0	7	0	8	4	0	0	5	0	6
Trypsin-1	P07477	TRY1_HUMAN	90	87	85	121	106	105	97	88	116	28	103	102	96	85	110	25	91	71	90	103	120	36	87	63	78	309	149	37
Trypsin-2	P07478	TRY2_HUMAN	89	96	106	139	135	88	69	77	118	44	140	102	84	82	128	40	110	138	54	162	106	65	99	126	62	347	137	63
Tubulin alpha-1B chain	P68363	TBA1B_HUMAN	1	1	4	3	0	0	0	0	5	1	0	0	0	0	5	1	0	0	5	7	9	0	0	0	5	5	8	0
Tubulin beta chain	P07437	TBB5_HUMAN	1	1	0	0	0	0	0	0	6	0	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0
Tubulin beta-2C chain	P68371	TBB2C_HUMAN	1	0	0	0	0	0	0	0	6	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
UDP-glucuronosyltransferase 2A3	Q6UWM9	UD2A3_HUMAN	1	1	0	0	0	0	0	1	4	0	0	0	0	1	3	0	0	0	0	0	0	0	0	0	0	0	0	0
UDP-glucuronosyltransferase 2B17	O75795	UDB17_HUMAN	0	0	0	0	0	0	0	1	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
UPF0451 protein C17orf61	Q8N2U0	CQ061_HUMAN	0	1	0	0	1	0	0	0	0	0	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
UPF0762 protein C6orf58	Q6P5S2	CF058_HUMAN	14	14	5	7	12	5	18	18	11	18	13	4	19	19	7	22	5	0	12	14	0	0	8	0	21	12	0	0
Uteroglobin	P11684	UTER_HUMAN	1	1	0	0	0	0	0	0	0	7	0	0	0	0	0	8	0	0	0	0	0	0	0	0	0	0	0	0
Villin-1	P09327	VILI_HUMAN	6	6	0	0	0	3	12	4	13	1	0	3	13	5	13	1	0	0	0	0	0	0	0	0	0	0	0	0
Vitamin D-binding protein	P02774	VTDB_HUMAN	14	13	0	0	1	13	15	11	12	32	1	13	14	11	10	31	0	0	0	0	0	0	0	0	0	0	0	0
Vitronectin	P04004	VTNC_HUMAN	8	10	1	1	0	9	14	6	5	12	0	12	17	7	6	16	5	0	0	0	0	0	4	0	0	0	0	0
Voltage-dependent anion-selective channel protein 1	P21796	VDAC1_HUMAN	1	1	0	0	0	0	0	2	5	0	0	0	0	2	5	0	0	0	0	0	0	0	0	0	0	0	0	0
WD repeat-containing protein 1	O75083	WDR1_HUMAN	2	2	0	0	1	1	1	2	5	0	2	1	1	2	5	0	0	0	0	0	0	0	0	0	0	0	0	0
Xaa-Pro aminopeptidase 2	O43895	XPP2_HUMAN	1	1	3	3	2	0	0	1	0	5	2	0	0	1	0	5	0	11	0	0	0	8	0	9	0	0	0	9
Xaa-Pro dipeptidase	P12955	PEPD_HUMAN	4	4	5	5	4	3	4	5	4	2	4	3	4	6	3	2	5	5	5	7	9	0	4	5	5	5	8	0
Xanthine dehydrogenase/oxidase	P47989	XDH_HUMAN	1	1	7	6	2	2	1	1	0	0	2	2	1	1	0	0	10	11	0	7	9	3	12	9	0	5	8	3
Zinc-alpha-2-glycoprotein	P25311	ZA2G_HUMAN	18	16	4	3	12	14	12	10	12	50	10	11	9	7	10	47	5	0	5	0	0	11	4	0	5	0	0	9
Zymogen granule membrane protein 16	O60844	ZG16_HUMAN	2	2	2	2	1	2	3	5	2	1	1	2	3	5	2	1	0	0	0	14	0	0	0	0	0	12	0	0
Zymogen granule protein 16 homolog B	Q96DA0	ZG16B_HUMAN	17	17	9	8	0	0	0	0	0	99	0	0	0	0	0	101	19	0	0	0	0	36	17	0	0	0	0	30