

Chymotrypsin-C	Q99895	0	0	0	0	0	0	0	13	0	0	0	4	0	13	0	1	1
Chymotrypsin-like elastase family member 2A	P08217	0	0	0	17	0	30	0	13	0	0	16	4	0	47	13	0	2
Chymotrypsin-like elastase family member 3A	P09093	0	0	9	0	14	32	0	0	0	3	15	0	9	46	0	1	
Chymotrypsin-like elastase family member 3B	P08861	17	0	0	8	0	0	0	20	0	6	3	7	17	8	20	1	
Clathrin heavy chain 1	Q00610	54	0	0	78	0	0	47	78	0	18	26	42	54	78	125	1	
Clathrin light chain A	P09496	0	14	0	0	0	0	0	0	0	5	0	0	14	0	0	1	
Coagulation factor XIII A chain	P00488	0	0	0	25	0	0	0	0	0	0	8	0	0	25	0	0	
Cofilin-1	P23528	42	78	53	36	34	30	47	0	14	58	33	20	173	100	61	3	
Collagen alpha-1	P02452	0	103	0	0	0	144	0	0	105	34	48	35	103	144	105	1	
Collagen alpha-1	P12109	89	91	0	65	96	68	0	67	56	60	76	41	180	229	123	2	
Collagen alpha-1	P39060	0	54	100	0	43	84	38	81	68	51	43	62	154	127	187	2	
Collagen alpha-1	Q05707	0	56	82	0	84	73	0	65	70	46	52	45	138	157	135	2	
Collagen alpha-1	Q99715	0	123	0	143	0	0	0	154	197	41	48	117	123	143	351	1	
Collagen alpha-2	P08123	0	0	0	0	0	173	0	193	201	0	58	131	0	173	394	0	
Collagen alpha-2	P12110	139	131	176	93	137	98	74	137	108	149	109	106	446	328	319	3	
Collagen alpha-3	P12111	218	260	355	213	393	242	172	226	232	278	283	210	833	848	630	3	
Complement C3	P01024	82	87	0	0	96	103	0	0	0	56	66	0	169	199	0	2	
Complement C4-A	P0COL4	0	0	0	0	0	0	0	0	56	0	0	19	0	0	56	0	
Complement C4-B	P0COL5	0	0	0	0	0	80	0	0	0	0	27	0	0	80	0	0	
Complement component C9	P02748	0	0	0	0	0	0	0	0	21	0	0	7	0	0	21	0	
Coronin-1A	P31146	15	0	0	0	0	0	0	0	0	5	0	0	15	0	0	1	
Coronin-1C	Q9ULV4	0	0	41	0	0	0	0	0	0	14	0	0	41	0	0	1	
C-type mannose receptor 2	Q9UBG0	0	0	0	0	0	34	0	0	26	0	11	9	0	34	26	0	
Cysteine and glycine-rich protein 1	P21291	7	0	21	0	14	0	0	0	16	9	5	5	28	14	16	2	
Cytoskeleton-associated protein 4	Q07065	54	0	0	40	0	46	34	0	0	18	29	11	54	86	34	1	
Cytosol aminopeptidase	P28838	0	0	0	27	0	0	0	0	0	0	9	0	0	27	0	0	
D-3-phosphoglycerate dehydrogenase	O43175	0	0	0	25	0	30	0	40	0	0	18	13	0	55	40	0	
Decorin	P07585	47	58	0	38	79	0	0	0	0	35	39	0	105	117	0	2	
Deleted in malignant brain tumors 1 protein	Q9UGM3	72	76	79	0	0	73	63	63	73	76	24	66	227	73	199	3	
Desmin	P17661	109	0	197	0	0	0	67	0	0	102	0	22	306	0	67	2	
Dihydropyrimidinase-related protein 2	Q16555	40	20	0	0	41	0	0	0	47	20	14	16	60	41	47	2	
Dolichyl-diphosphooligosaccharide-- glycosyltransferase 1	P04843	0	0	0	0	0	0	0	47	0	0	0	16	0	0	47	0	
ELAV-like protein 1	Q15717	0	0	0	0	0	0	24	0	0	0	0	8	0	24	0	0	
Elongation factor 1-gamma	P26641	0	0	0	23	0	0	20	31	0	0	8	17	0	23	51	0	
Elongation factor 2	P13639	0	0	0	40	0	0	0	51	0	0	13	17	0	40	51	0	
EMILIN-1	Q9Y6C2	69	131	91	44	101	64	45	58	101	97	70	68	291	209	204	3	
Endoplasmic	P14625	0	0	0	36	0	0	33	0	0	0	12	11	0	36	33	0	
Eosinophil cationic protein	P12724	49	14	0	0	0	18	22	7	26	21	6	18	63	18	55	2	
Eosinophil peroxidase	P11678	99	0	0	0	0	0	0	0	0	33	0	0	99	0	0	1	
Epiplakin	P58107	0	0	0	0	0	0	165	0	171	0	0	112	0	0	336	0	
Erythrocyte band 7 integral membrane protein	P27105	0	0	0	0	0	0	0	0	28	0	0	9	0	0	28	0	
F-actin-capping protein subunit alpha-1	P52907	0	0	0	32	0	0	0	0	0	0	11	0	0	32	0	0	
F-actin-capping protein subunit alpha-2	P47755	0	0	0	23	0	0	0	0	0	0	8	0	0	23	0	0	
Fatty acid synthase	P49327	0	0	0	0	0	0	45	0	0	0	0	15	0	0	45	0	
Fatty acid-binding protein, liver	P07148	0	0	0	0	0	0	13	0	0	0	0	4	0	13	0	0	
Ferritin heavy chain	P02794	0	0	0	6	31	0	0	0	23	0	12	8	0	37	23	0	
Ferritin light chain	P02792	0	22	18	11	5	14	11	11	26	13	10	16	40	30	48	2	
Fibrillin-1	P35555	0	56	147	53	43	0	0	0	70	68	32	23	203	96	70	2	
Fibrillin-2	P35556	0	70	0	0	0	0	0	49	0	23	0	16	70	0	49	1	
Fibrinogen alpha chain	P02671	0	64	0	0	0	98	0	0	0	21	33	0	64	98	0	1	
Fibrinogen beta chain	P02675	0	58	0	0	0	62	0	0	0	19	21	0	58	62	0	1	
Fibronectin	P02751	0	87	0	93	0	66	38	0	138	29	53	59	87	159	176	1	
Fibulin-1	P23142	20	16	23	27	22	11	20	0	16	20	20	12	59	60	36	3	
Filamin-A	P21333	237	217	560	171	247	162	209	215	295	338	193	240	1014	580	719	3	
Filamin-B	O75369	0	0	0	0	0	0	51	0	0	0	0	17	0	0	51	0	
Fructose-bisphosphate aldolase A	P04075	32	58	23	36	34	34	62	36	35	38	35	44	113	104	133	3	
Gamma-glutamyltransferase 5	P36269	0	0	0	0	17	0	0	0	28	0	6	9	0	17	28	0	
Gelsolin	P06396	0	0	65	42	0	0	0	0	0	22	14	0	65	42	0	1	
Glucagon	P01275	47	54	26	40	172	30	0	34	77	42	81	37	127	242	111	3	
Glucose-6-phosphate isomerase	P06744	0	0	32	32	0	0	22	0	0	11	11	7	32	32	22	1	
Glucosidase 2 subunit beta	P14314	0	0	0	48	0	0	0	0	0	0	16	0	0	48	0	0	

Ig kappa chain V-IV region B17	P06314	0	10	9	11	0	0	0	0	6	4	0	19	11	0	2	1	0
Ig lambda chain V-I region NEWM	P01703	0	14	15	13	0	9	0	0	0	7	0	29	22	0	2	2	0
IgG Fc-binding protein	Q9Y6R7	0	0	0	0	0	0	42	0	0	0	14	0	0	42	0	0	1
Immunoglobulin J chain	P01591	15	8	0	2	0	11	5	9	14	8	4	9	23	13	28	2	3
Importin subunit beta-1	Q14974	0	0	0	0	0	0	22	0	0	0	0	7	0	22	0	1	
Inner centromere protein	Q9NQ57	67	0	0	0	0	0	0	0	22	0	0	67	0	0	1	0	
Insulin	P01308	0	0	0	0	22	0	0	0	0	7	0	0	22	0	0	1	0
Integrin alpha-M	P11215	0	0	0	0	0	0	0	0	68	0	0	23	0	68	0	0	1
Junction plakoglobin	P14923	0	0	0	0	43	0	0	0	0	14	0	0	43	0	0	1	0
Keratin, type I cuticular Ha3-I	O76009	20	0	0	0	0	0	0	0	7	0	0	20	0	1	0	0	
Keratin, type I cuticular Ha6	O76013	0	0	0	0	0	0	31	0	0	0	10	0	31	0	0	1	0
Keratin, type I cytoskeletal 10	P13645	272	133	232	116	196	132	149	150	129	212	148	143	637	444	428	3	3
Keratin, type I cytoskeletal 14	P02533	79	52	79	67	89	43	60	72	112	70	66	81	210	199	244	3	3
Keratin, type I cytoskeletal 16	P08779	0	0	0	0	0	0	53	0	0	0	0	18	0	53	0	1	0
Keratin, type I cytoskeletal 18	P05783	0	0	0	0	38	50	62	40	33	0	29	45	0	88	135	0	3
Keratin, type I cytoskeletal 19	P08727	45	36	79	0	0	52	63	51	66	53	17	60	160	52	180	3	3
Keratin, type I cytoskeletal 9	P35527	240	99	235	179	278	189	141	146	173	191	215	153	574	646	460	3	3
Keratin, type II cuticular Hb5	P78386	0	0	0	0	0	0	33	0	0	0	0	11	0	33	0	1	0
Keratin, type II cuticular Hb6	O43790	42	0	0	0	0	0	0	0	0	14	0	0	42	0	1	0	0
Keratin, type II cytoskeletal 1	P04264	331	187	379	196	295	237	190	240	300	299	243	243	897	728	730	3	3
Keratin, type II cytoskeletal 2 epidermal	P35908	257	109	238	109	175	125	120	121	112	201	136	118	604	409	353	3	3
Keratin, type II cytoskeletal 5	P13647	153	0	0	0	0	0	69	0	0	51	0	23	153	0	69	1	1
Keratin, type II cytoskeletal 7	P08729	0	0	0	0	0	0	0	60	0	0	0	20	0	60	0	0	1
Keratin, type II cytoskeletal 8	P05787	92	38	0	0	0	59	114	60	87	43	20	87	130	59	261	2	3
Lactotransferrin	O2788	0	0	0	0	0	0	22	0	37	0	0	20	0	59	0	0	2
Laminin subunit alpha-5	O15230	0	99	0	0	0	0	0	0	0	33	0	0	99	0	1	0	0
Lithostathine-1-beta	P48304	0	0	0	0	0	5	0	0	0	0	2	0	5	0	0	1	0
L-lactate dehydrogenase A chain	P00338	0	0	0	0	0	0	25	0	30	0	0	18	0	55	0	0	2
Lumican	P51884	0	34	0	25	36	7	9	18	42	11	23	23	34	68	69	1	3
Lysozyme C	P61626	0	0	0	27	0	0	22	0	28	0	9	17	0	50	0	1	2
Lysyl oxidase homolog 1	Q08397	0	0	0	0	48	0	0	0	0	0	16	0	48	0	0	1	0
Macrophage migration inhibitory factor	P14174	0	22	0	0	0	0	24	27	0	7	0	17	22	0	51	1	2
Macrophage-capping protein	P40121	0	28	0	27	0	0	0	11	19	9	9	10	28	27	30	1	2
Malate dehydrogenase, mitochondrial	P40926	0	12	0	17	19	11	0	22	0	4	16	7	12	47	22	1	1
Matrin-3	P43243	45	0	0	42	0	0	0	0	0	15	14	0	45	42	0	1	0
Mimecan	P20774	0	0	0	0	36	0	0	0	0	0	12	0	36	0	0	1	0
Moesin	P26038	0	0	0	29	0	0	0	0	0	0	10	0	29	0	0	1	0
Mucin-2	Q02817	0	0	0	0	0	0	34	0	0	0	0	11	0	34	0	0	1
Mucin-5B	Q9HC84	0	0	0	0	0	0	0	0	91	0	0	30	0	91	0	0	1
Myeloblastin	P24158	0	0	0	0	0	0	18	0	9	0	0	9	0	27	0	0	2
Myeloperoxidase	P05164	92	95	0	72	113	75	109	96	220	62	87	142	187	260	425	2	3
Myosin light polypeptide 6	P60660	0	18	41	19	24	0	15	0	12	20	14	9	59	43	27	2	2
Myosin-11	P35749	0	0	185	0	0	0	0	0	0	62	0	0	185	0	1	0	0
Myosin-14	Q72406	0	0	0	0	0	0	123	0	0	0	0	41	0	123	0	0	1
Neurosecretory protein VGF	O15240	0	0	0	0	31	0	0	0	0	10	0	0	31	0	0	1	0
Neutrophil elastase	P08246	40	10	0	0	17	25	53	0	80	17	14	44	50	42	133	2	2
Nicotinate phosphoribosyltransferase	Q6XQN6	0	0	0	0	0	0	7	0	0	0	0	2	0	7	0	0	1
Nidogen-1	P14543	0	50	53	0	26	0	18	31	63	34	9	37	103	26	112	2	3
Non-histone chromosomal protein HMG-17	P05204	0	0	0	29	0	0	13	0	0	0	10	4	0	29	13	0	1
Non-POU domain-containing octamer-binding protein	Q15233	32	0	0	0	0	0	31	0	0	11	0	10	32	0	31	1	1
Non-secretory ribonuclease	P10153	0	0	0	0	0	7	0	0	0	0	2	0	7	0	0	1	0
Outer dense fiber protein 3	Q96PU9	0	0	0	0	0	0	0	0	14	0	0	5	0	14	0	0	1
Pancreatic alpha-amylase	P04746	0	0	0	0	29	0	0	0	0	10	0	0	29	0	0	1	0
Pancreatic prohormone	P01298	40	0	0	0	0	0	0	0	26	13	0	9	40	0	26	1	1
Pancreatic triacylglycerol lipase	P16233	0	0	0	0	0	27	0	0	0	0	9	0	27	0	0	1	0
PDZ and LIM domain protein 1	O00151	0	0	0	0	0	0	18	0	0	0	0	6	0	18	0	0	1
Peptidyl-prolyl cis-trans isomerase A	P62937	47	44	35	53	29	50	42	72	33	42	44	49	126	132	147	3	3
Peptidyl-prolyl cis-trans isomerase FKBP11	Q9NYL4	0	0	0	0	0	5	0	20	0	0	2	7	0	20	0	1	1
Periostin	Q15063	0	66	0	46	34	0	0	0	56	22	27	19	66	80	56	1	1
Peroxiredoxin-1	Q06830	20	30	26	25	0	0	0	18	0	25	8	6	76	25	18	3	1
Peroxiredoxin-4	Q13162	0	0	0	21	10	23	9	40	0	0	18	16	0	54	49	0	2

Tripeptidyl-peptidase 1	O14773	0	0	0	0	0	0	11	0	19	0	0	10	0	30	0	0	2	
Tropomyosin alpha-1 chain	P09493	0	0	67	0	29	0	0	0	22	10	0	67	29	0	1	1	0	
Tropomyosin alpha-4 chain	P67936	20	0	0	29	0	14	20	20	30	7	14	23	20	43	70	1	2	3
Tropomyosin beta chain	P07951	0	0	88	0	0	0	0	0	0	29	0	0	88	0	0	1	0	0
Trypsin-1	P07477	116	0	0	82	0	80	47	92	0	39	54	46	116	162	139	1	2	2
Trypsin-2	P07478	0	0	0	69	67	73	0	69	0	0	70	23	0	209	69	0	3	1
Trypsin-3	P35030	0	0	0	0	0	0	0	63	0	0	0	21	0	0	63	0	0	1
Tryptase alpha/beta-1	Q15661	22	0	26	17	22	32	0	36	49	16	24	28	48	71	85	2	3	2
Tryptophan--tRNA ligase, cytoplasmic	P23381	0	0	0	13	0	0	0	0	0	0	4	0	0	13	0	0	1	0
Tubulin alpha-1A chain	Q71U36	0	0	0	0	38	0	0	0	0	0	13	0	0	38	0	0	1	0
Tubulin alpha-1B chain	P68363	0	0	32	0	0	0	0	0	0	11	0	0	32	0	0	1	0	0
Tubulin alpha-1C chain	Q9BQE3	0	14	0	23	0	34	40	16	0	5	19	19	14	57	56	1	2	2
Tubulin beta chain	P07437	37	54	32	19	41	41	49	0	54	41	34	34	123	101	103	3	3	2
Tubulin beta-4B chain	P68371	0	0	0	0	0	0	45	25	0	0	0	23	0	0	70	0	0	2
Tubulin-specific chaperone A	O75347	0	0	0	0	0	0	9	0	0	0	0	3	0	9	0	0	0	1
Ubiquitin-40S ribosomal protein S27a	P62979	62	48	59	44	0	55	0	40	0	56	33	13	169	99	40	3	2	1
Ubiquitin-60S ribosomal protein L40	P62987	0	0	0	0	38	0	40	0	0	0	13	13	0	38	40	0	1	1
Ubiquitin-like modifier-activating enzyme 1	P22314	45	0	0	0	0	0	0	0	0	15	0	0	45	0	0	1	0	0
Versican core protein	P13611	84	109	0	57	69	0	0	74	73	64	42	49	193	126	147	2	2	2
Vimentin	P08670	104	135	109	78	134	96	74	103	108	116	103	95	348	308	285	3	3	3
Vinculin	P18206	82	68	111	107	105	84	82	101	0	87	99	61	261	296	183	3	3	2
Vitronectin	P04004	0	18	0	0	0	0	0	0	40	6	0	13	18	0	40	1	0	1
V-type proton ATPase subunit B, brain isoform	P21281	0	0	0	23	0	0	11	0	0	0	8	4	0	23	11	0	1	1

AIP: autoimmune pancreatitis; CP: chronic pancreatitis; PC: pancreatic cancer

Supplementary Table 2. Twenty-four proteins identified in both autoimmune pancreatitis and chronic pancreatitis specimens and not in pancreatic cancer specimens, as illustrated in the Venn diagram in Figure 2.

Protein names	UniProt ID	Spectral counts						Mean spectral counts		Total spectral counts		No. of specimens	
		AIP#1	AIP#2	AIP#3	CP#1	CP#2	CP#3	AIP	CP	AIP	CP	AIP	CP
60S ribosomal protein L23	P62829	35	0	0	42	0	36	12	26	35	78	1	2
60S ribosomal protein L26	P61254	20	0	0	29	26	39	7	31	20	94	1	3
60S ribosomal protein L34	P49207	0	28	0	27	31	0	9	19	28	58	1	2
Asporin	Q9BXN1	7	0	0	0	41	0	2	14	7	41	1	1
Chymotrypsin-like elastase family member 3A	P09093	0	0	9	0	14	32	3	15	9	46	1	2
Complement C3	P01024	82	87	0	0	96	103	56	66	169	199	2	2
Decorin	P07585	47	58	0	38	79	0	35	39	105	117	2	2
Fibrinogen alpha chain	P02671	0	64	0	0	0	98	21	33	64	98	1	1
Fibrinogen beta chain	P02675	0	58	0	0	0	62	19	21	58	62	1	1
Gelsolin	P06396	0	0	65	42	0	0	22	14	65	42	1	1
Heat shock 70 kDa protein 1A/1B	P08107	0	0	59	0	36	0	20	12	59	36	1	1
Hemoglobin subunit delta	P02042	0	137	0	69	0	0	46	23	137	69	1	1
Histone H2A.V	Q71UI9	32	40	0	27	0	0	24	9	72	27	2	1
Histone H2B type 1-M	Q99879	0	155	0	185	120	0	52	102	155	305	1	2
Ig kappa chain V-III region VG	P04433	0	10	0	11	0	0	3	4	10	11	1	1
Ig kappa chain V-IV region B17	P06314	0	10	9	11	0	0	6	4	19	11	2	1
Ig lambda chain V-I region NEWM	P01703	0	14	15	13	0	9	10	7	29	22	2	2
Matrin-3	P43243	45	0	0	42	0	0	15	14	45	42	1	1
Plasma cell-induced resident endoplasmic reticulum protein	Q8WU39	25	36	15	34	0	0	25	11	76	34	3	1
Probable phosphoglycerate mutase 4	Q8N0Y7	0	14	0	0	26	0	5	9	14	26	1	1
Prolargin	P51888	0	40	0	21	0	0	13	7	40	21	1	1
T-complex protein 1 subunit beta	P78371	25	0	0	25	0	0	8	8	25	25	1	1
Thrombospondin-1	P07996	0	52	0	78	0	48	17	42	52	126	1	2
Tropomyosin alpha-1 chain	P09493	0	0	67	0	29	0	22	10	67	29	1	1

AIP: autoimmune pancreatitis; CP: chronic pancreatitis

Supplementary Table 3. Nineteen proteins identified in both autoimmune pancreatitis and pancreatic cancer specimens and not in chronic pancreatitis specimens, as illustrated in the Venn diagram in Figure 2.

Protein names	UniProt ID	Spectral counts						Mean spectral counts		Total spectral counts		No. of specimens	
		AIP#1	AIP#2	AIP#3	PC#1	PC#2	PC#3	AIP	PC	AIP	PC	AIP	PC
40S ribosomal protein S12	P25398	0	10	0	0	27	0	3	9	10	27	1	1
Actin, aortic smooth muscle	P62736	0	0	493	294	0	0	164	98	493	294	1	1
Alpha-actinin-1	P12814	0	0	79	0	0	37	26	12	79	37	1	1
Caldesmon	Q05682	0	0	147	51	0	0	49	17	147	51	1	1
Desmin	P17661	109	0	197	67	0	0	102	22	306	67	2	1
Fibrillin-2	P35556	0	70	0	0	49	0	23	16	70	49	1	1
Heterogeneous nuclear ribonucleoprotein U	Q00839	104	0	0	87	0	0	35	29	104	87	1	1
Histone H2A type 1-C	Q93077	0	97	0	0	87	75	32	54	97	162	1	2
Histone H2B type 1-D	P58876	0	0	185	0	119	0	62	40	185	119	1	1
Keratin, type II cytoskeletal 5	P13647	153	0	0	69	0	0	51	23	153	69	1	1
Macrophage migration inhibitory factor	P14174	0	22	0	24	27	0	7	17	22	51	1	2
Non-POU domain-containing octamer-binding protein	Q15233	32	0	0	31	0	0	11	10	32	31	1	1
Pancreatic prohormone	P01298	40	0	0	0	0	26	13	9	40	26	1	1
Plastin-2	P13796	45	54	44	0	45	0	48	15	143	45	3	1
Protein S100-A6	P06703	0	6	0	7	0	7	2	5	6	14	1	2
Splicing factor, proline- and glutamine-rich	P23246	0	56	0	54	0	63	19	39	56	117	1	2
Succinate dehydrogenase	P31040	27	0	44	20	0	0	24	7	71	20	2	1
Transitional endoplasmic reticulum ATPase	P55072	0	54	0	51	49	0	18	33	54	100	1	2
Vitronectin	P04004	0	18	0	0	0	40	6	13	18	40	1	1

AIP: autoimmune pancreatitis; PC: pancreatic cancer

Supplementary Table 4. Forty-nine proteins identified in both chronic pancreatitis and pancreatic cancer specimens and not in the autoimmune pancreatitis specimens, as illustrated in the Venn diagram in Figure 2.

Protein names	UniProt ID	Spectral counts						Mean spectral counts		Total spectral counts		No. of specimens	
		CP#1	CP#2	CP#3	PC#1	PC#2	PC#3	CP	PC	CP	PC	CP	PC
14-3-3 protein gamma	P61981	0	0	32	34	0	33	11	22	32	67	1	2
14-3-3 protein theta	P27348	17	12	0	16	0	12	10	9	29	28	2	2
40S ribosomal protein S10	P46783	2	0	0	9	18	0	1	9	2	27	1	2
40S ribosomal protein S14	P62263	0	0	36	29	20	0	12	16	36	49	1	2
40S ribosomal protein S15a	P62244	19	0	32	20	0	0	17	7	51	20	2	1
40S ribosomal protein S28	P62857	23	0	0	0	16	0	8	5	23	16	1	1
40S ribosomal protein S4, X isoform	P62701	0	0	27	0	47	0	9	16	27	47	1	1
40S ribosomal protein S6	P62753	38	43	57	38	0	0	46	13	138	38	3	1
60S ribosomal protein L14	P50914	36	0	39	0	63	0	25	21	75	63	2	1
60S ribosomal protein L29	P47914	55	0	34	0	60	0	30	20	89	60	2	1
60S ribosomal protein L35	P42766	21	0	0	24	0	0	7	8	21	24	1	1
Alpha-amylase 2B	P19961	32	0	0	0	25	0	11	8	32	25	1	1
ATP-citrate synthase	P53396	0	57	0	38	0	0	19	13	57	38	1	1
Brain acid soluble protein 1	P80723	6	0	0	0	0	16	2	5	6	16	1	1
Carboxypeptidase B	P15086	0	0	21	0	22	0	7	7	21	22	1	1
Cartilage oligomeric matrix protein	P49747	21	0	0	0	0	26	7	9	21	26	1	1
Chymotrypsin-like elastase family member 2A	P08217	17	0	30	0	13	0	16	4	47	13	2	1
Collagen alpha-2	P08123	0	0	173	0	193	201	58	131	173	394	1	2
C-type mannose receptor 2	Q9UBG0	0	0	34	0	0	26	11	9	34	26	1	1
D-3-phosphoglycerate dehydrogenase	O43175	25	0	30	0	40	0	18	13	55	40	2	1
Elongation factor 1-gamma	P26641	23	0	0	20	31	0	8	17	23	51	1	2
Elongation factor 2	P13639	40	0	0	0	51	0	13	17	40	51	1	1
Endoplasmin	P14625	36	0	0	33	0	0	12	11	36	33	1	1
Ferritin heavy chain	P02794	6	31	0	0	0	23	12	8	37	23	2	1
Gamma-glutamyltransferase 5	P36269	0	17	0	0	0	28	6	9	17	28	1	1
Heat shock cognate 71 kDa protein	P11142	27	0	0	58	0	0	9	19	27	58	1	1
Histone H1.5	P16401	78	0	0	0	0	84	26	28	78	84	1	1
Histone H2A type 2-C	Q16777	67	0	112	85	0	0	60	28	179	85	2	1
Histone H2B type 1-J	P06899	0	98	0	129	0	0	33	43	98	129	1	1
Histone H3.2	Q71DI3	126	0	0	85	0	105	42	63	126	190	1	2
Ig kappa chain V-III region HIC	P18136	21	0	0	2	0	0	7	1	21	2	1	1
Keratin, type I cytoskeletal 18	P05783	0	38	50	62	40	33	29	45	88	135	2	3
Lysozyme C	P61626	27	0	0	22	0	28	9	17	27	50	1	2
Non-histone chromosomal protein HMG-17	P05204	29	0	0	13	0	0	10	4	29	13	1	1
Peptidyl-prolyl cis-trans isomerase FKBP11	Q9NYL4	0	0	5	0	20	0	2	7	5	20	1	1
Peroxiredoxin-4	Q13162	21	10	23	9	40	0	18	16	54	49	3	2
Phosphatidylethanolamine-binding protein 1	P30086	0	7	0	0	9	0	2	3	7	9	1	1
Plectin	Q15149	0	242	0	0	0	265	81	88	242	265	1	1
Protein CutA	O60888	8	12	0	7	0	0	7	2	20	7	2	1
Protein disulfide-isomerase A2	Q13087	0	0	36	0	49	0	12	16	36	49	1	1
Protein S100-A9	P06702	0	14	0	11	0	63	5	25	14	74	1	2
Ribosome-binding protein 1	Q9P2E9	0	0	91	0	154	0	30	51	91	154	1	1
Serpin H1	P50454	25	0	27	25	0	30	17	18	52	55	2	2

T-complex protein 1 subunit epsilon	P48643	42	46	0	38	29	0	29	22	88	67	2	2
Tenascin	P24821	63	0	0	49	0	0	21	16	63	49	1	1
Transketolase	P29401	0	60	0	47	40	40	20	42	60	127	1	3
Trypsin-2	P07478	69	67	73	0	69	0	70	23	209	69	3	1
Ubiquitin-60S ribosomal protein L40	P62987	0	38	0	40	0	0	13	13	38	40	1	1
V-type proton ATPase subunit B, brain isoform	P21281	23	0	0	11	0	0	8	4	23	11	1	1

CP: chronic pancreatitis; PC: pancreatic cancer