

#	Identified Proteins (211)	Accession Number	ANOVA Test (p-value): (p < 0.05)	Quantitative Profile	Taxonomy	CS F-N-1A	CS F-N-1B	CS F-N-2A	CS F-N-2B	CSF -N-PM 1A	CSF -N-PM 1B	CSF -N-PM 2A	CSF -N-PM 2B	CSF -S-PM 1A	CSF -S-PM 1B	CSF -S-PM 2A	CSF -S-PM 2B
1	Apolipoprotein A-II OS=Homo sapiens GN=APOA2 PE=1 SV=1	APOA2_HUMAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	45	48	42	42	2	1	1	1	0	0	5	3
2	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4_HUMAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	40	29	31	26	0	1	2	0	0	0	0	0
3	Selenium-binding protein 1 OS=Homo sapiens GN=SELENBP1 PE=1 SV=2	SBP1_HUMAN	< 0,00010	CSF-N low, CSF-N-PM high, CSF-S-PM low	Homo sapiens	0	0	0	0	6	6	3	3	1	1	2	1
4	Tubulin beta-4B chain OS=Homo sapiens GN=TUBB4B PE=1 SV=1	TBB4B_HUMAN	< 0,00010	CSF-N low, CSF-N-PM high, CSF-S-PM low	Homo sapiens	0	0	0	1	11	13	14	9	0	0	1	0
5	Alpha-1B-glycoprotein OS=Homo sapiens GN=A1BG PE=1 SV=4	A1BG_HUMAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	33	40	33	32	0	0	4	3	3	2	11	12
6	Haptoglobin OS=Homo sapiens GN=HP PE=1 SV=1	HPT_HUMAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	16 0	16 0	90	85	0	0	0	0	5	1	24	0
7	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1	IGHG1_HUMAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	40 1	37 9	33 2	30 6	51	55	67	78	57	51	141	0
8	Astrocytic phosphoprotein	PEA15_HUMAN	< 0,00010	CSF-N low,	Homo	0	0	0	0	7	6	5	3	3	0	0	1

	PEA-15 OS=Homo sapiens GN=PEA15 PE=1 SV=2	UMAN		CSF-N-PM high, CSF-S- PM low	sapien s												
9	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3	A2MG_H UMAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapien s	30 8	28 4	13 8	14 9	0	0	1	2	10	10	7	0
10	Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1 SV=3	APOH_HU MAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapien s	12	11	17	17	0	0	1	0	1	2	5	0
11	Leucine-rich alpha-2- glycoprotein OS=Homo sapiens GN=LRG1 PE=1 SV=2	A2GL_HU MAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapien s	6	8	7	6	0	0	2	1	0	1	0	0
12	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	CLUS_HU MAN	< 0,00010	CSF-N high, CSF-N-PM high, CSF-S- PM low	Homo sapien s	83	57	73	77	41	39	41	49	5	3	12	0
13	Glyceraldehyde-3- phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	G3P_HU MAN	< 0,00010	CSF-N low, CSF-N-PM high, CSF-S- PM low	Homo sapien s	0	0	3	5	42	43	72	77	24	23	44	10
14	Ig gamma-4 chain C region OS=Homo sapiens GN=IGHG4 PE=1 SV=1	IGHG4_H UMAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapien s	27 8	22 6	22 8	21 7	41	52	47	45	42	42	114	0
15	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	CO3_HU MAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapien s	15 7	15 5	23 1	22 9	0	2	2	2	13	13	13	0
16	Apolipoprotein D OS=Homo sapiens	APOD_HU MAN	< 0,00010	CSF-N low, CSF-N-PM	Homo sapien	7	4	6	0	20	22	15	18	0	0	0	1

	GN=APOD PE=1 SV=1			high, CSF-S-PM low	s												
17	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	ITIH1_HUMAN	< 0,00010	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	11	11	8	15	0	0	0	0	1	2	1	0
18	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1	APOA1_HUMAN	0.00013	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	307	302	179	180	63	81	48	41	103	83	75	80
19	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	ITIH2_HUMAN	0.0002	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	2	5	5	4	0	0	0	0	0	0	0	0
20	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	ENOA_HUMAN	0.00023	CSF-N low, CSF-N-PM high, CSF-S-PM low	Homo sapiens	0	0	0	0	87	97	110	97	28	24	97	4
21	Ig gamma-3 chain C region OS=Homo sapiens GN=IGHG3 PE=1 SV=2	IGHG3_HUMAN	0.00024	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	241	176	226	208	36	44	41	44	44	41	107	0
22	14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1 SV=1	1433T_HUMAN	0.00028	CSF-N low, CSF-N-PM low, CSF-S-PM high	Homo sapiens	0	0	0	0	0	0	2	2	8	8	14	15
23	Ig kappa chain C region OS=Homo sapiens GN=IGKC PE=1 SV=1	IGKC_HUMAN	0.00037	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	291	310	471	446	41	39	58	50	21	20	81	2
24	Ig mu chain C region OS=Homo sapiens GN=IGHM PE=1 SV=3	IGHM_HUMAN	0.00044	CSF-N high, CSF-N-PM low, CSF-S-	Homo sapiens	46	47	15	19	0	0	0	0	0	0	0	0

				PM low													
25	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	HRG_HUMAN	0.00059	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	4	3	6	2	0	0	0	0	0	0	0	0
26	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA_HUMAN	0.00082	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	1	2	3	2	0	0	0	0	0	0	0	0
27	Hemopexin OS=Homo sapiens GN=HPX PE=1 SV=2	HEMO_HUMAN	0.00083	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	50	64	29	39	3	3	13	16	6	5	23	0
28	Ig gamma-2 chain C region OS=Homo sapiens GN=IGHG2 PE=1 SV=2	IGHG2_HUMAN	0.00085	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	19 6	20 9	14 1	14 6	44	51	31	26	43	43	106	0
29	Apolipoprotein C-III OS=Homo sapiens GN=APOC3 PE=1 SV=1	APOC3_HUMAN	0.00091	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	15	11	6	7	0	0	0	0	0	0	0	4
30	Dickkopf-related protein 3 OS=Homo sapiens GN=DKK3 PE=1 SV=2	DKK3_HUMAN	0.0015	CSF-N low, CSF-N-PM high, CSF-S- PM low	Homo sapiens	0	0	7	3	7	5	11	10	0	0	1	0
31	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2	ANXA5_HUMAN	0.0015	CSF-N low, CSF-N-PM high, CSF-S- PM low	Homo sapiens	0	0	0	0	2	2	3	6	0	1	1	0
32	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	VTNC_HUMAN	0.002	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	12	15	26	23	0	2	0	0	0	1	7	0

33	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	CO4B_HUMAN	0.0022	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	52	57	108	116	1	2	1	2	17	10	8	0
34	Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1 PE=1 SV=2	IGHA1_HUMAN	0.003	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	45	52	43	33	2	1	29	20	2	2	20	0
35	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFAH_HUMAN	0.0031	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	4	4	6	10	0	0	0	0	0	0	0	0
36	Alpha-2-HS-glycoprotein OS=Homo sapiens GN=AHSG PE=1 SV=1	FETUA_HUMAN	0.0034	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	7	15	9	5	0	2	0	1	1	2	0	1
37	Immunoglobulin heavy variable 3-53 OS=Homo sapiens GN=IGHV3-53 PE=1 SV=2	HV353_HUMAN	0.004	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapiens	9	9	4	6	0	0	0	2	1	0	5	0
38	Neural cell adhesion molecule 1 OS=Homo sapiens GN=NCAM1 PE=1 SV=3	NCAM1_HUMAN	0.0051	CSF-N low, CSF-N-PM high, CSF-S- PM low	Homo sapiens	0	0	1	2	4	3	1	4	0	0	0	0
39	Phosphatidylethanolamine- binding protein 1 OS=Homo sapiens GN=PEBP1 PE=1 SV=3	PEBP1_HUMAN	0.0057	CSF-N low, CSF-N-PM high, CSF-S- PM low	Homo sapiens	0	1	0	0	22	22	33	35	16	21	30	9
40	Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5	PRDX2_HUMAN	0.0069	CSF-N low, CSF-N-PM high, CSF-S- PM low	Homo sapiens	0	0	6	8	32	29	15	12	7	7	3	23
41	Chromogranin-A OS=Homo	CMGA_H	0.0081	CSF-N low,	Homo	0	0	0	1	4	7	11	10	0	0	0	0

	sapiens GN=CHGA PE=1 SV=7	UMAN		CSF-N-PM high, CSF-S-PM low	sapiens												
42	Amyloid-like protein 1 OS=Homo sapiens GN=APLP1 PE=1 SV=3	APLP1_HUMAN	0.0083	CSF-N low, CSF-N-PM high, CSF-S-PM low	Homo sapiens	0	0	0	0	2	1	4	2	0	0	0	0
43	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1	1433E_HUMAN	0.0085	CSF-N low, CSF-N-PM high, CSF-S-PM high	Homo sapiens	0	0	3	0	13	16	19	17	10	4	24	3
44	Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2 PE=1 SV=1	LAC2_HUMAN	0.0096	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	66	66	89	103	12	7	64	53	15	14	0	0
45	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB_HUMAN	0.0098	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	12	10	39	25	0	0	0	0	0	0	0	0
46	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	CO5_HUMAN	0.012	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	6	4	1	1	0	0	0	0	0	0	0	0
47	Heat shock 70 kDa protein 1B OS=Homo sapiens GN=HSPA1B PE=1 SV=1	HS71B_HUMAN (+1)	0.013	CSF-N low, CSF-N-PM high, CSF-S-PM low	Homo sapiens	0	0	0	0	2	2	7	6	4	0	1	0
48	Actin, cytoplasmic 2 OS=Homo sapiens GN=ACTG1 PE=1 SV=1	ACTG_HUMAN	0.013	CSF-N low, CSF-N-PM high, CSF-S-PM low	Homo sapiens	0	0	22	15	60	61	16	24	16	7	17	2
49	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLMN_HUMAN	0.013	CSF-N high, CSF-N-PM	Homo sapiens	2	2	6	7	0	0	0	0	0	0	1	0

				low, CSF-S-PM low	s												
50	Immunoglobulin J chain OS=Homo sapiens GN=JCHAIN PE=1 SV=4	IGJ_HUMAN	0.014	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	0	2	2	1	0	0	0	0	0	0	0	0
51	Beta-2-microglobulin OS=Homo sapiens GN=B2M PE=1 SV=1	B2MG_HUMAN	0.014	CSF-N high, CSF-N-PM high, CSF-S-PM low	Homo sapiens	2	3	5	3	2	2	2	4	0	0	0	0
52	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE_HUMAN	0.015	CSF-N high, CSF-N-PM high, CSF-S-PM low	Homo sapiens	2	1	40	44	33	25	57	56	9	5	10	0
53	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFAB_HUMAN	0.016	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	4	4	10	12	0	0	0	0	1	2	4	0
54	Immunoglobulin kappa variable 3-20 OS=Homo sapiens GN=IGKV3-20 PE=1 SV=2	KV320_HUMAN	0.018	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	13	9	6	5	0	0	0	1	6	8	0	0
55	Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1	TBA1B_HUMAN	0.02	CSF-N low, CSF-N-PM high, CSF-S-PM low	Homo sapiens	0	0	0	2	21	20	12	11	0	0	19	0
56	Fructose-bisphosphate aldolase C OS=Homo sapiens GN=ALDOC PE=1 SV=2	ALDOC_HUMAN	0.022	CSF-N low, CSF-N-PM low, CSF-S-PM high	Homo sapiens	0	0	0	4	4	2	7	12	48	35	31	1
57	Gamma-enolase OS=Homo sapiens GN=ENO2 PE=1 SV=3	ENOG_HUMAN	0.023	CSF-N low, CSF-N-PM high, CSF-S-	Homo sapiens	0	0	0	0	47	54	31	21	10	7	40	9

				PM low													
58	Alpha-1-acid glycoprotein 1 OS=Homo sapiens GN=ORM1 PE=1 SV=1	A1AG1_H UMAN	0.026	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapien s	52	69	69	76	37	37	31	42	37	34	60	22
59	14-3-3 protein eta OS=Homo sapiens GN=YWHAH PE=1 SV=4	1433F_H UMAN	0.032	CSF-N low, CSF-N-PM low, CSF-S- PM high	Homo sapien s	0	0	0	1	0	0	1	0	9	3	8	0
60	Serum amyloid A-1 protein OS=Homo sapiens GN=SAA1 PE=1 SV=1	SAA1_HU MAN	0.036	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapien s	6	5	49	45	0	0	0	0	0	0	0	0
61	Triosephosphate isomerase OS=Homo sapiens GN=TPI1 PE=1 SV=3	TPIS_HU MAN	0.038	CSF-N low, CSF-N-PM high, CSF-S- PM low	Homo sapien s	0	0	1	3	10	16	35	42	8	3	15	25
62	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2	HBB_HU MAN	0.038	CSF-N low, CSF-N-PM low, CSF-S- PM high	Homo sapien s	18	21	91	94	113 8	116 8	27	24	163 4	174 5	175	170 6
63	Secretogranin-1 OS=Homo sapiens GN=CHGB PE=1 SV=2	SCG1_HU MAN	0.038	CSF-N low, CSF-N-PM high, CSF-S- PM low	Homo sapien s	0	0	0	0	2	2	8	4	0	0	0	0
64	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	KNG1_HU MAN	0.039	CSF-N high, CSF-N-PM low, CSF-S- PM low	Homo sapien s	2	2	6	8	0	0	0	0	2	0	2	0
65	Flavin reductase (NADPH) OS=Homo sapiens GN=BLVRB PE=1 SV=3	BLVRB_H UMAN	0.039	CSF-N low, CSF-N-PM high, CSF-S- PM low	Homo sapien s	0	0	0	0	4	3	0	1	1	0	0	1

66	Prostaglandin-H2 D-isomerase OS=Homo sapiens GN=PTGDS PE=1 SV=1	PTGDS_HUMAN	0.04	CSF-N low, CSF-N-PM high, CSF-S-PM low	Homo sapiens	9	14	31	21	33	33	66	80	20	19	34	30
67	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	VTDB_HUMAN	0.045	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	15	13	59	59	19	20	0	2	1	1	5	1
68	Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1 SV=3	A1AT_HUMAN	0.045	CSF-N high, CSF-N-PM high, CSF-S-PM low	Homo sapiens	376	380	237	251	90	92	239	264	85	81	148	85
69	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	ANT3_HUMAN	0.049	CSF-N high, CSF-N-PM low, CSF-S-PM low	Homo sapiens	6	2	29	27	0	0	0	0	0	0	4	0
70	Hemoglobin subunit delta OS=Homo sapiens GN=HBD PE=1 SV=2	HBD_HUMAN	0.05	[]	Homo sapiens	0	0	31	26	663	676	11	0	917	934	42	916
71	Cystatin-C OS=Homo sapiens GN=CST3 PE=1 SV=1	CYTC_HUMAN	0.052	[]	Homo sapiens	15	12	22	27	33	40	96	131	5	6	0	6
72	Hemoglobin subunit gamma-2 OS=Homo sapiens GN=HBG2 PE=1 SV=2	HBG2_HUMAN	0.053	[]	Homo sapiens	0	5	7	8	146	177	4	3	245	238	4	156
73	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1	GELS_HUMAN	0.054	[]	Homo sapiens	3	4	5	7	4	2	16	15	3	0	2	0
74	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	IC1_HUMAN	0.054	[]	Homo sapiens	14	6	33	38	0	0	15	13	0	2	1	0
75	Rab GDP dissociation	GDIA_HUMAN	0.054	[]	Homo	0	0	1	1	0	0	2	1	6	4	10	0

	inhibitor alpha OS=Homo sapiens GN=GDI1 PE=1 SV=2	MAN			sapiens												
76	Neurosecretory protein VGF OS=Homo sapiens GN=VGF PE=1 SV=2	VGF_HU MAN	0.056	[]	Homo sapiens	0	0	0	0	3	2	17	11	0	0	0	0
77	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	CO9_HU MAN	0.058	[]	Homo sapiens	5	7	1	0	0	0	0	0	0	0	0	0
78	Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3	TRFE_HU MAN	0.059	[]	Homo sapiens	29 5	32 2	26 5	24 8	183	197	281	254	105	110	322	9
79	Retinol-binding protein 4 OS=Homo sapiens GN=RBP4 PE=1 SV=3	RET4_HU MAN	0.061	[]	Homo sapiens	7	8	0	1	0	0	0	0	0	0	0	0
80	Carbonic anhydrase 2 OS=Homo sapiens GN=CA2 PE=1 SV=2	CAH2_HU MAN	0.062	[]	Homo sapiens	0	0	0	0	19	19	86	93	18	18	75	37
81	Carbonic anhydrase 1 OS=Homo sapiens GN=CA1 PE=1 SV=2	CAH1_HU MAN	0.068	[]	Homo sapiens	0	0	1	1	29	27	0	0	28	24	24	14
82	Creatine kinase B-type OS=Homo sapiens GN=CKB PE=1 SV=1	KCRB_HU MAN	0.069	[]	Homo sapiens	0	0	0	0	14	17	76	77	42	35	112	63
83	Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1 SV=2	FIBB_HU MAN	0.07	[]	Homo sapiens	20	17	57	52	28	24	0	0	3	5	8	0
84	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	AACT_HU MAN	0.073	[]	Homo sapiens	62	53	10 8	10 4	3	4	59	52	26	26	51	0
85	Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4	FINC_HU MAN	0.075	[]	Homo sapiens	2	1	31	31	0	0	0	0	0	0	0	0

86	L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2	LDHB_HUMAN	0.082	[]	Homo sapiens	0	0	4	7	14	12	38	35	15	13	45	0
87	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4	EZRI_HUMAN	0.084	[]	Homo sapiens	0	0	1	0	6	7	16	24	1	3	22	0
88	Thy-1 membrane glycoprotein OS=Homo sapiens GN=THY1 PE=1 SV=2	THY1_HUMAN	0.086	[]	Homo sapiens	0	0	0	0	1	0	3	1	0	0	1	0
89	Fatty acid-binding protein, epidermal OS=Homo sapiens GN=FABP5 PE=1 SV=3	FABP5_HUMAN	0.086	[]	Homo sapiens	0	0	0	0	2	1	13	9	1	2	5	7
90	Extracellular superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD3 PE=1 SV=2	SODE_HUMAN	0.087	[]	Homo sapiens	2	1	0	0	1	0	9	6	0	0	0	0
91	Carbonyl reductase [NADPH] 1 OS=Homo sapiens GN=CBR1 PE=1 SV=3	CBR1_HUMAN	0.089	[]	Homo sapiens	0	0	0	0	5	7	23	21	6	8	28	1
92	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3	PGK1_HUMAN	0.089	[]	Homo sapiens	0	0	1	0	11	9	28	22	3	5	34	0
93	ProSAAS OS=Homo sapiens GN=PCSK1N PE=1 SV=1	PCSK1_HUMAN	0.093	[]	Homo sapiens	0	0	0	0	0	2	17	14	3	2	0	0
94	Alpha-1-acid glycoprotein 2 OS=Homo sapiens GN=ORM2 PE=1 SV=2	A1AG2_HUMAN	0.094	[]	Homo sapiens	28	33	24	30	12	0	18	0	20	23	36	0
95	Glutathione S-transferase Mu 3 OS=Homo sapiens	GSTM3_HUMAN	0.096	[]	Homo sapiens	0	0	0	0	0	0	7	13	12	9	16	0

	GN=GSTM3 PE=1 SV=3				s												
96	Fibrinogen alpha chain OS=Homo sapiens GN=FGA PE=1 SV=2	FIBA_HUMAN	0.097	[]	Homo sapiens	2	6	21	24	6	10	1	3	1	0	2	1
97	Retinal dehydrogenase 1 OS=Homo sapiens GN=ALDH1A1 PE=1 SV=2	AL1A1_HUMAN	0.1	[]	Homo sapiens	0	0	0	0	0	0	0	0	8	8	0	0
98	Osteopontin OS=Homo sapiens GN=SPP1 PE=1 SV=1	OSTP_HUMAN	0.1	[]	Homo sapiens	6	4	3	3	0	2	5	6	0	0	0	0
99	Spectrin alpha chain, non-erythrocytic 1 OS=Homo sapiens GN=SPTAN1 PE=1 SV=3	SPTN1_HUMAN	0.1	[]	Homo sapiens	0	0	0	0	0	0	0	0	6	0	5	0
100	Platelet glycoprotein Ib beta chain OS=Homo sapiens GN=GP1BB PE=1 SV=1	GP1BB_HUMAN	0.1	[]	Homo sapiens	0	0	0	0	4	3	0	0	0	0	0	0
101	Ig delta chain C region OS=Homo sapiens GN=IGHD PE=1 SV=2	IGHD_HUMAN	0.1	[]	Homo sapiens	0	0	6	7	0	0	0	0	0	0	0	0
102	Coagulation factor XIII A chain OS=Homo sapiens GN=F13A1 PE=1 SV=4	F13A_HUMAN	0.11	[]	Homo sapiens	0	0	0	0	3	4	0	0	0	0	0	0
103	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R_HUMAN	0.11	[]	Homo sapiens	0	0	4	3	0	0	0	0	0	0	0	0
104	Insulin-like growth factor-binding protein complex acid labile subunit OS=Homo sapiens GN=IGFALS PE=1 SV=1	ALS_HUMAN	0.11	[]	Homo sapiens	0	0	4	3	0	0	0	0	0	0	0	0

105	Tubulin alpha-4A chain OS=Homo sapiens GN=TUBA4A PE=1 SV=1	TBA4A_HUMAN	0.11	[]	Homo sapiens	0	0	0	1	22	21	0	3	0	0	13	0
106	Tubulin beta-1 chain OS=Homo sapiens GN=TUBB1 PE=1 SV=1	TBB1_HUMAN	0.11	[]	Homo sapiens	0	0	1	1	23	22	0	0	0	0	0	0
107	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4	MYH9_HUMAN	0.11	[]	Homo sapiens	0	0	4	4	98	113	0	0	0	0	0	0
108	Integrin beta-3 OS=Homo sapiens GN=ITGB3 PE=1 SV=2	ITB3_HUMAN	0.11	[]	Homo sapiens	0	0	0	0	8	12	0	0	0	0	0	0
109	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4	PEDF_HUMAN	0.11	[]	Homo sapiens	20	18	18	24	3	4	31	30	1	3	13	0
110	Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2	HBA_HUMAN	0.12	[]	Homo sapiens	2	4	54	46	1529	1530	12	7	1325	1296	42	1458
111	Sorcin OS=Homo sapiens GN=SRI PE=1 SV=1	SORCN_HUMAN	0.12	[]	Homo sapiens	0	0	0	0	0	0	5	4	0	0	0	1
112	IgGfc-binding protein OS=Homo sapiens GN=FCGBP PE=1 SV=3	FCGBP_HUMAN	0.12	[]	Homo sapiens	0	0	4	6	0	0	0	0	0	0	0	0
113	Lipopolysaccharide-binding protein OS=Homo sapiens GN=LBP PE=1 SV=3	LBP_HUMAN	0.12	[]	Homo sapiens	0	0	4	6	0	0	0	0	0	0	0	0
114	Filamin-A OS=Homo sapiens GN=FLNA PE=1 SV=4	FLNA_HUMAN	0.12	[]	Homo sapiens	0	0	3	2	43	32	0	0	0	0	0	0
115	Immunoglobulin lambda-like polypeptide 5	IGLL5_HUMAN	0.12	[]	Homo sapiens	54	57	57	69	9	6	62	51	15	14	45	6

	OS=Homo sapiens GN=IGLL5 PE=2 SV=2				s												
116	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	LUM_HUMAN	0.13	[]	Homo sapiens	0	1	0	0	0	0	7	9	1	0	0	0
117	Alpha-actinin-1 OS=Homo sapiens GN=ACTN1 PE=1 SV=2	ACTN1_HUMAN	0.13	[]	Homo sapiens	0	0	5	3	19	14	1	0	0	0	0	0
118	Vinculin OS=Homo sapiens GN=VCL PE=1 SV=4	VINC_HUMAN	0.13	[]	Homo sapiens	0	0	2	3	15	14	0	0	0	0	0	0
119	Acyl-CoA-binding protein OS=Homo sapiens GN=DBI PE=1 SV=2	ACBP_HUMAN	0.13	[]	Homo sapiens	0	0	0	0	0	0	3	5	0	0	0	0
120	Protein FAM3C OS=Homo sapiens GN=FAM3C PE=1 SV=1	FAM3C_HUMAN	0.13	[]	Homo sapiens	0	0	0	0	0	0	9	12	0	0	0	3
121	Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3	COF1_HUMAN	0.14	[]	Homo sapiens	0	0	0	0	2	1	12	8	10	8	5	28
122	Na(+)/H(+) exchange regulatory cofactor NHERF1 OS=Homo sapiens GN=SLC9A3R1 PE=1 SV=4	NHRF1_HUMAN	0.14	[]	Homo sapiens	0	0	0	0	0	0	10	8	0	0	0	4
123	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	THRB_HUMAN	0.14	[]	Homo sapiens	0	0	2	1	0	0	0	0	0	0	0	0
124	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	CO6_HUMAN	0.14	[]	Homo sapiens	0	0	6	3	0	0	0	0	0	0	0	0
125	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2	GSTP1_HUMAN	0.14	[]	Homo sapiens	0	0	0	0	5	3	71	63	14	11	8	44

126	Cellular retinoic acid-binding protein 1 OS=Homo sapiens GN=CRABP1 PE=1 SV=2	RABP1_H UMAN	0.14	[]	Homo sapiens	0	0	0	0	10	8	2	0	5	8	1	2
127	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4	VIME_HU MAN	0.14	[]	Homo sapiens	0	0	0	1	66	64	6	6	90	82	27	0
128	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2	ALDOA_H UMAN	0.14	[]	Homo sapiens	0	0	0	0	1	1	10	7	3	3	13	19
129	Glutathione S-transferase Mu 2 OS=Homo sapiens GN=GSTM2 PE=1 SV=2	GSTM2_H UMAN	0.14	[]	Homo sapiens	0	0	0	0	0	0	5	5	7	6	2	20
130	Beta-1,4-glucuronyltransferase 1 OS=Homo sapiens GN=B4GAT1 PE=1 SV=1	B4GA1_H UMAN	0.14	[]	Homo sapiens	0	0	0	0	0	0	10	13	0	0	0	4
131	Protein ABHD14B OS=Homo sapiens GN=ABHD14B PE=1 SV=1	ABHEB_H UMAN	0.14	[]	Homo sapiens	0	0	0	0	0	0	2	2	0	0	0	1
132	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S_HUM AN	0.15	[]	Homo sapiens	0	0	11	5	0	0	0	0	0	0	0	0
133	Protein deglycase DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2	PARK7_H UMAN	0.15	[]	Homo sapiens	0	0	0	0	2	0	21	19	11	5	6	16
134	Peroxiredoxin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3	PRDX6_H UMAN	0.15	[]	Homo sapiens	0	0	2	0	1	1	30	27	4	3	4	18
135	Erythrocyte band 7 integral membrane protein OS=Homo sapiens	STOM_H UMAN	0.15	[]	Homo sapiens	0	0	3	2	17	12	0	0	0	0	0	0

	GN=STOM PE=1 SV=3																
136	Ubiquitin-conjugating enzyme E2 N OS=Homo sapiens GN=UBE2N PE=1 SV=1	UBE2N_HUMAN	0.15	[]	Homo sapiens	0	0	0	0	0	0	1	2	0	0	0	0
137	Peptidyl-prolyl cis-trans isomerase FKBP1A OS=Homo sapiens GN=FKBP1A PE=1 SV=2	FKB1A_HUMAN	0.15	[]	Homo sapiens	0	0	0	0	0	0	1	2	0	0	0	0
138	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1	HSP7C_HUMAN	0.16	[]	Homo sapiens	0	0	1	0	6	3	14	19	8	10	32	0
139	Protein S100-B OS=Homo sapiens GN=S100B PE=1 SV=2	S100B_HUMAN	0.16	[]	Homo sapiens	0	0	0	0	0	0	17	21	2	4	1	7
140	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	CO7_HUMAN	0.16	[]	Homo sapiens	0	0	5	2	0	0	0	0	0	0	0	0
141	Phospholipid transfer protein OS=Homo sapiens GN=PLTP PE=1 SV=1	PLTP_HUMAN	0.16	[]	Homo sapiens	0	0	5	2	0	0	0	0	0	0	0	0
142	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2	PPIA_HUMAN	0.16	[]	Homo sapiens	0	0	0	0	12	8	57	49	27	27	6	87
143	Talin-1 OS=Homo sapiens GN=TLN1 PE=1 SV=3	TLN1_HUMAN	0.16	[]	Homo sapiens	0	0	31	23	87	79	0	0	0	0	0	0
144	Zinc-alpha-2-glycoprotein OS=Homo sapiens GN=AZGP1 PE=1 SV=2	ZA2G_HUMAN	0.16	[]	Homo sapiens	9	10	2	0	0	0	3	2	0	0	2	0
145	Apolipoprotein A-IV OS=Homo sapiens	APOA4_HUMAN	0.16	[]	Homo sapiens	30	32	1	0	0	0	4	4	1	1	0	3

	GN=APOA4 PE=1 SV=3				s												
146	Fibrinogen gamma chain OS=Homo sapiens GN=FGG PE=1 SV=3	FIBG_HUMAN	0.17	[]	Homo sapiens	8	5	25	22	11	9	0	0	6	1	8	0
147	Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5 PE=1 SV=4	PRDX5_HUMAN	0.17	[]	Homo sapiens	0	0	0	0	2	5	50	44	10	9	15	49
148	Ras-related protein Rap-1b OS=Homo sapiens GN=RAP1B PE=1 SV=1	RAP1B_HUMAN	0.17	[]	Homo sapiens	0	0	5	4	24	16	0	0	0	0	0	0
149	Adenylate kinase isoenzyme 1 OS=Homo sapiens GN=AK1 PE=1 SV=3	KAD1_HUMAN	0.17	[]	Homo sapiens	0	0	0	0	4	5	26	23	4	8	33	0
150	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1	PRDX1_HUMAN	0.17	[]	Homo sapiens	0	0	0	0	12	10	25	27	10	6	14	68
151	Cystatin-B OS=Homo sapiens GN=CSTB PE=1 SV=2	CYTB_HUMAN	0.17	[]	Homo sapiens	0	0	0	0	1	2	12	3	0	1	0	3
152	Macrophage migration inhibitory factor OS=Homo sapiens GN=MIF PE=1 SV=4	MIF_HUMAN	0.17	[]	Homo sapiens	0	0	0	0	3	3	9	9	5	5	2	21
153	Calmodulin OS=Homo sapiens GN=CALM1 PE=1 SV=2	CALM_HUMAN	0.17	[]	Homo sapiens	0	0	0	0	0	0	2	3	1	0	0	1
154	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2	PROF1_HUMAN	0.18	[]	Homo sapiens	0	0	0	0	6	5	12	14	1	0	0	27
155	Tropomyosin alpha-3 chain OS=Homo sapiens GN=TPM3 PE=1 SV=2	TPM3_HUMAN	0.18	[]	Homo sapiens	0	0	0	0	3	1	0	0	0	0	0	0

156	Inositol-3-phosphate synthase 1 OS=Homo sapiens GN=ISYNA1 PE=1 SV=1	INO1_HUMAN	0.18	[]	Homo sapiens	0	0	0	0	0	0	4	3	0	0	3	0
157	Phosphoglycerate mutase 1 OS=Homo sapiens GN=PGAM1 PE=1 SV=2	PGAM1_HUMAN	0.18	[]	Homo sapiens	0	0	0	0	0	0	23	17	2	3	10	3
158	Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4	G6PI_HUMAN	0.18	[]	Homo sapiens	0	0	0	0	0	0	17	12	8	8	14	0
159	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1	1433Z_HUMAN	0.18	[]	Homo sapiens	0	0	9	5	17	21	5	6	11	12	25	4
160	Integrin alpha-IIb OS=Homo sapiens GN=ITGA2B PE=1 SV=3	ITA2B_HUMAN	0.18	[]	Homo sapiens	0	0	2	6	14	11	0	0	0	0	0	0
161	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	AMBP_HUMAN	0.19	[]	Homo sapiens	16	19	0	0	0	0	0	0	0	0	8	0
162	Chloride intracellular channel protein 6 OS=Homo sapiens GN=CLIC6 PE=2 SV=3	CLIC6_HUMAN	0.2	[]	Homo sapiens	0	0	0	0	0	0	12	12	0	1	6	0
163	Superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD1 PE=1 SV=2	SODC_HUMAN	0.21	[]	Homo sapiens	0	0	0	0	1	0	3	3	4	5	19	0
164	Transketolase OS=Homo sapiens GN=TKT PE=1 SV=3	TKT_HUMAN	0.23	[]	Homo sapiens	0	0	0	0	3	1	4	0	2	3	11	0
165	Chitinase-3-like protein 1 OS=Homo sapiens GN=CHI3L1 PE=1 SV=2	CH3L1_HUMAN	0.23	[]	Homo sapiens	9	2	0	0	0	0	0	0	0	0	0	0
166	Monocyte differentiation	CD14_HUMAN	0.23	[]	Homo	0	0	13	3	0	0	0	0	0	0	0	0

	antigen CD14 OS=Homo sapiens GN=CD14 PE=1 SV=2	MAN			sapiens												
167	Thrombospondin-1 OS=Homo sapiens GN=THBS1 PE=1 SV=2	TSP1_HUMAN	0.24	[]	Homo sapiens	0	0	16	11	21	20	0	0	0	0	0	0
168	Inositol monophosphatase 1 OS=Homo sapiens GN=IMPA1 PE=1 SV=1	IMPA1_HUMAN	0.24	[]	Homo sapiens	0	0	0	0	0	0	1	4	0	0	0	0
169	Ubiquitin carboxyl-terminal hydrolase isozyme L1 OS=Homo sapiens GN=UCHL1 PE=1 SV=2	UCHL1_HUMAN	0.25	[]	Homo sapiens	0	0	0	0	0	0	2	1	1	2	4	22
170	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	ALBU_HUMAN	0.26	[]	Homo sapiens	3056	2905	2149	2079	673	658	2196	2066	1319	1444	3263	396
171	Protein-L-isoaspartate(D-aspartate) O-methyltransferase OS=Homo sapiens GN=PCMT1 PE=1 SV=4	PIMT_HUMAN	0.27	[]	Homo sapiens	0	0	0	0	0	0	1	2	3	1	0	8
172	Beta-Ala-His dipeptidase OS=Homo sapiens GN=CNDP1 PE=1 SV=4	CNDP1_HUMAN	0.28	[]	Homo sapiens	2	0	2	2	1	2	7	7	2	6	7	1
173	Glial fibrillary acidic protein OS=Homo sapiens GN=GFAP PE=1 SV=1	GFAP_HUMAN	0.29	[]	Homo sapiens	90	91	0	0	2	1	5	7	40	28	105	1
174	Dihydropyrimidinase-related protein 2 OS=Homo sapiens GN=DPYSL2 PE=1 SV=1	DPYL2_HUMAN	0.3	[]	Homo sapiens	0	0	0	1	2	1	11	10	6	4	29	0
175	Pyruvate kinase PKM OS=Homo sapiens GN=PKM	KPYM_HUMAN	0.3	[]	Homo sapiens	0	0	4	7	4	4	60	54	17	16	73	0

	PE=1 SV=4				s												
176	Fermitin family homolog 3 OS=Homo sapiens GN=FERMT3 PE=1 SV=1	URP2_HUMAN	0.3	[]	Homo sapiens	0	0	2	4	3	8	0	0	0	0	0	0
177	Transaldolase OS=Homo sapiens GN=TALDO1 PE=1 SV=2	TALDO_HUMAN	0.32	[]	Homo sapiens	0	0	0	0	0	0	0	0	1	0	9	0
178	Secernin-1 OS=Homo sapiens GN=SCRN1 PE=1 SV=2	SCRN1_HUMAN	0.32	[]	Homo sapiens	0	0	0	0	0	0	1	1	1	0	3	0
179	Malate dehydrogenase, cytoplasmic OS=Homo sapiens GN=MDH1 PE=1 SV=4	MDHC_HUMAN	0.33	[]	Homo sapiens	0	0	0	0	1	0	2	4	3	1	19	0
180	GTP-binding nuclear protein Ran OS=Homo sapiens GN=RAN PE=1 SV=3	RAN_HUMAN	0.36	[]	Homo sapiens	0	0	0	0	0	0	2	1	1	1	0	6
181	Synaptic vesicle membrane protein VAT-1 homolog-like OS=Homo sapiens GN=VAT1L PE=1 SV=2	VAT1L_HUMAN	0.36	[]	Homo sapiens	0	0	0	0	0	0	11	9	3	3	22	0
182	Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	ANGT_HUMAN	0.38	[]	Homo sapiens	20	10	74	58	10	9	66	54	19	32	11	18
183	14-3-3 protein gamma OS=Homo sapiens GN=YWHAG PE=1 SV=2	1433G_HUMAN	0.38	[]	Homo sapiens	0	0	3	0	0	6	2	2	4	2	11	0
184	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CERU_HUMAN	0.39	[]	Homo sapiens	26	16	39	34	16	18	7	15	11	9	39	0
185	Phosphoserine aminotransferase	SERC_HUMAN	0.4	[]	Homo sapiens	0	0	0	0	0	0	1	3	0	1	4	0

	OS=Homo sapiens GN=PSAT1 PE=1 SV=2				s												
186	Tubulin polymerization-promoting protein OS=Homo sapiens GN=TPPP PE=1 SV=1	TPPP_HUMAN	0.41	[]	Homo sapiens	0	0	0	0	0	0	0	0	0	0	14	0
187	Lactoylglutathione lyase OS=Homo sapiens GN=GLO1 PE=1 SV=4	LGUL_HUMAN	0.41	[]	Homo sapiens	0	0	0	0	0	0	4	0	0	0	0	0
188	S-formylglutathione hydrolase OS=Homo sapiens GN=ESD PE=1 SV=2	ESTD_HUMAN	0.41	[]	Homo sapiens	0	0	0	0	0	0	0	0	0	0	0	4
189	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	CO8G_HUMAN	0.41	[]	Homo sapiens	0	0	0	2	0	0	0	0	0	0	0	0
190	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	C1QC_HUMAN	0.41	[]	Homo sapiens	0	2	0	0	0	0	0	0	0	0	0	0
191	Calcyphosin OS=Homo sapiens GN=CAPS PE=1 SV=1	CAYP1_HUMAN	0.41	[]	Homo sapiens	0	0	0	0	0	0	2	0	0	1	45	
192	N(G),N(G)-dimethylarginine dimethylaminohydrolase 1 OS=Homo sapiens GN=DDAH1 PE=1 SV=3	DDAH1_HUMAN	0.42	[]	Homo sapiens	0	0	0	0	0	0	1	2	0	0	5	1
193	Low molecular weight phosphotyrosine protein phosphatase OS=Homo sapiens GN=ACP1 PE=1 SV=3	PPAC_HUMAN	0.43	[]	Homo sapiens	0	0	0	0	0	0	1	4	0	0	0	4
194	Retinol-binding protein 1 OS=Homo sapiens	RET1_HUMAN	0.44	[]	Homo sapiens	0	0	0	0	0	0	11	13	0	0	0	25

	GN=RBP1 PE=1 SV=2				s													
195	Calbindin OS=Homo sapiens GN=CALB1 PE=1 SV=2	CALB1_HUMAN	0.44	[]	Homo sapiens	0	0	0	0	0	0	0	0	1	1	2	0	0
196	Aflatoxin B1 aldehyde reductase member 2 OS=Homo sapiens GN=AKR7A2 PE=1 SV=3	ARK72_HUMAN	0.46	[]	Homo sapiens	0	0	0	0	0	0	2	0	0	0	0	0	16
197	Cytosolic non-specific dipeptidase OS=Homo sapiens GN=CNDP2 PE=1 SV=2	CNDP2_HUMAN	0.46	[]	Homo sapiens	0	0	0	0	0	0	1	0	0	0	0	6	0
198	ADP-ribosylation factor 1 OS=Homo sapiens GN=ARF1 PE=1 SV=2	ARF1_HUMAN (+1)	0.48	[]	Homo sapiens	0	0	0	0	0	0	2	2	1	0	0	0	13
199	14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB PE=1 SV=3	1433B_HUMAN	0.49	[]	Homo sapiens	0	0	3	0	0	6	1	2	4	1	9	0	0
200	Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3	TAGL2_HUMAN	0.5	[]	Homo sapiens	0	0	0	0	4	3	1	0	0	0	0	0	10
201	Fatty acid-binding protein, adipocyte OS=Homo sapiens GN=FABP4 PE=1 SV=3	FABP4_HUMAN	0.5	[]	Homo sapiens	0	0	0	0	0	0	6	4	0	0	0	9	0
202	Galectin-1 OS=Homo sapiens GN=LGALS1 PE=1 SV=2	LEG1_HUMAN	0.5	[]	Homo sapiens	0	0	0	0	0	0	2	4	0	0	0	0	9
203	Keratin, type II cytoskeletal 1 OS=Homo sapiens GN=KRT1 PE=1 SV=6	K2C1_HUMAN	0.51	[]	Homo sapiens	2	0	2	1	3	0	4	2	3	1	14	0	0
204	F-box only protein 2 OS=Homo sapiens	FBX2_HUMAN	0.52	[]	Homo sapiens	0	0	0	0	0	0	2	1	0	0	0	7	0

	GN=FBXO2 PE=1 SV=2				s												
205	D-dopachrome decarboxylase OS=Homo sapiens GN=DDT PE=1 SV=3	DOPD_HUMAN	0.52	[]	Homo sapiens	0	0	0	0	0	0	1	3	0	0	0	6
206	Ribonuclease inhibitor OS=Homo sapiens GN=RNH1 PE=1 SV=2	RINI_HUMAN	0.53	[]	Homo sapiens	0	0	0	0	0	0	3	0	0	0	1	0
207	Aspartate aminotransferase, cytoplasmic OS=Homo sapiens GN=GOT1 PE=1 SV=3	AATC_HUMAN	0.54	[]	Homo sapiens	0	0	0	0	0	0	2	1	0	0	5	0
208	Nucleoside diphosphate kinase B OS=Homo sapiens GN=NME2 PE=1 SV=1	NDKB_HUMAN	0.54	[]	Homo sapiens	0	0	0	0	0	0	1	3	0	0	7	0
209	Transthyretin OS=Homo sapiens GN=TTR PE=1 SV=1	TTHY_HUMAN	0.55	[]	Homo sapiens	319	316	61	69	51	60	210	202	51	44	98	219
210	Ig heavy chain V-III region NIE OS=Homo sapiens PE=1 SV=1	HV309_HUMAN	0.61	[]	Homo sapiens	9	3	0	0	0	0	2	2	2	0	2	0
211	Phosphoglucomutase-1 OS=Homo sapiens GN=PGM1 PE=1 SV=3	PGM1_HUMAN	0.72	[]	Homo sapiens	0	0	2	0	0	0	2	1	1	0	0	0