

Supplementary Material to “Extracellular vesicles in infectious diseases caused by protozoan parasites in buffaloes”

Additional file 3. Summary of extracellular vesicle proteins of *Theileria* spp. isolated from serum of treated animals present in Fraction F3.

Code access	Description*	MW [kDa]	Coverage	Unique peptides	Score
E1BNR0	Apolipoprotein	515.438	24.12962557	103	674,8379145
Q2UVX4	Complement C3	187.135	33.65442505	56	440,4326642
A0A0F6QNP7	Complement component 3	187.064	33.23299217	55	440,4326642
G3X7A5	Complement C3	187.028	32.69114991	55	410,7775776
B8Y9S9	Fibronectin 1	262.263	27.06325932	3	327,4399574
G5E5A9	Fibronectin	271.952	25.14124294	2	321,4439662
P07589	Fibronectin	271.983	25.14124294	42	321,4439662
B8Y9T0	Fibronectin 1	248.974	26.54320988	21	318,3170657
P15497	Apolipoprotein A-I	30.258	19.7564276	24	220,1763519
Q0VCM4	Glycogen phosphorylase	97.394	19.72972973	38	220,1763519
F1MI18	Uncharacterized protein	165.654	59.62264151	22	232,1050236
F1MJK3	Uncharacterized protein	165.506	8.790035587	0	115,8492713
A0A140T897	Serum albumin	69.278	8.796296296	19	115,8492713
P02769	Serum albumin	69.248	27.22371968	5	404,0782466
P23805	Conglutinin	37.971	26.1589404	3	873,9068512
P80457	Xanthine dehydrogenase/oxidase	146.696	14,44444444	14	65,70898986
F1MUT3	Xanthine dehydrogenase/oxidase	146.669	22,73476112	9	73,49145103
B0JYQ0	ALB protein	69.248	22,73476112	15	73,49145103
F1MVP0	ADAM metalloproteinase with thrombospondin type 1	151.374	48,55769231	13	119,6688337
F1N0R5	von Willebrand factor	307.475	15,57377049	0	100,6843615
F1MAV0	Fibrinogen beta chain	56.405	13,41880342	12	62,5247829
E1BH06	Uncharacterized protein	192.644	30,3030303	10	52,56129336
F5XVA9	von Willebrand factor	307.745	32,05128205	5	52,56129336

Code access	Description*	MW [kDa]	Coverage	Unique peptides	Score
P02676	Fibrinogen beta chain	53.306	20,59308072	6	62,91113496
F1MY85	Complement C5a anaphylatoxin	188.927	13,60655738	3	100,6843615
V6F869	Apolipoprotein A-I-like	23.629	44,6685879	12	108,6491276
A0A0F6QMJ3	Complement component 5	188.691	25,67324955	8	303,3408718
F1N3Q7	Apolipoprotein A-IV	42.963	8,737864078	0	71,69110668
Q32PJ2	Apolipoprotein A-IV	42.991	19,0070922	15	31,68743467
G3N0I4	Cytosol aminopeptidas	54.047	13,95749569	10	111,2448874
P00727	Cytosol aminopeptidas	56.254	26,31578947	3	33,63287568
Q693V9	Complement component 3d	34.422	26,31578947	10	33,63287568
Q3MHL4	Adenosylhomocysteinase 3	47.607	30,0330033	2	76,51224542
P00432	Catalase	59.878	12,44239631	9	25,84321988
Q7SIH1	Alpha-2-macroglobulin	167.47	24,29048414	3	646,0610726
Q95KV5	Fibronectin	38.32	6,357388316	5	20,59954548
F1MQ37	Myosin heavy chain 9	226.962	6,357388316	4	20,59954548
F1MD73	Uncharacterized protein	189.978	4,975347378	8	20,59954548
A5PJE3	Fibrinogen alpha chain	66.957	38,75	0	71,74341667
P02672	Fibrinogen alpha chain	66.971	14,77477477	5	62,25517595
A5D9E9	Complement C1r subcomponent precursor	80.161	13,49593496	9	39,98661864
Q03247	Apolipoprotein	35.958	13,49593496	8	39,98661864
Q2TBU0	Haptoglobin	44.831	27,84810127	0	25,54059851
E1BI98	Collagen type VI alpha 1 chain	108.603	27,84810127	4	25,54059851
G5E513	Uncharacterized protein	49.939	27,84810127	6	25,54059851
A6QPX7	FGB protein (Fragment)	37.907	28,48484848	8	30,96007919
P01030	Complement C4	101.817	28,18627451	9	267,9672889
E1B805	Uncharacterized protein	187.149	28,96725441	12	267,9672889
G3X6K8	Haptoglobin	44.845	24,31289641	15	267,9672889
A0A140T881	Apolipoprotein	36.017	10,21582734	7	32,74262905
G3N3D4	Potassium channel tetramerization domain containing	35.701	10,30478955	7	32,74262905
A7YWR0	Apolipoprotein	36.051	14,4469526	19	38,56606507
E1BFN6	Dihydropyrimidinase	56.267	14,71264368	20	38,56606507
P81187	Complement factor B	85.312	27,154047	0	227,8457786
P17697	Clusterin	51.081	17,30418944	13	42,44953275
F1N076	Ceruloplasmin	123.731	36,24454148	7	32,37882257

Code access	Description*	MW [kDa]	Coverage	Unique peptides	Score
R9QSM8	Alpha-2-macroglobulin	133.302	25,43520309	2	505,1839079
F1N514	CD5 antigen-like precursor	50.305	32,38512035	0	770,8895055
Q3SZZ9	FGG protein	49.136	22,92993631	12	60,16956151
Q0VCX1	Complement C1s subcomponent	76.56	5,501618123	9	11,86303473
F1MJ12	Complement C1s subcomponent	77.332	13,30798479	3	32,08415091
F1MGU7	Fibrinogen gamma-B chain	50.2	13,30798479	5	32,08415091
Q95KV4	Fibronectin (Fragment)	35.464	31,64556962	1	14,37422311
Q3Y5Z3	Adiponectin	26.117	11,03603604	0	33,17564273
Q0ZCB4	Apolipoprotein	27.057	27,27272727	3	148,6272044
A2VDY5	Hydroxysteroid (17-beta)	28.378	20,38834951	9	65,30266976
Q9MYP6	17-beta-hydroxysteroid dehydrogenas	28.404	25,41666667	19	22,7700392
A6QNW7	CD5L protein	50.179	10,93394077	5	23,2725662
Q0P5J7	Keratin. type I cuticular Ha5	49.905	12,14128035	21	87,20624363
B7FEK7	43kDa collectin	33.594	12,19512195	18	87,20624363
E1BPW6	Keratin. type I cuticular Ha5	49.739	10,54347826	7	51,87173843
Q28178	Thrombospondin-1	129.451	28,76106195	5	551,6399485
A7E3W2	Galectin-3-binding protein	62.087	8,333333333	10	14,14523625
F1MKG2	Collagen type VI alpha 2 chain	109.253	17,81376518	3	26,02097392
B8Y898	Malic enzyme	63.853	12,40875912	6	23,08411241
P02663	Alpha-S2-casein	26.002	7,755775578	7	24,12152457
F1MC11	Keratin. type I cytoskeletal 14	51.879	48,59813084	9	65,37755024
A6QNZ7	Keratin 10	54.816	5,176767677	10	9,596704483
P12799	Fibrinogen gamma-B chain	50.212	12,43654822	23	11,27226317
Q3SYR8	Immunoglobulin J chain	17.846	9,009009009	0	5,487135172
F1N3A1	Thrombospondin-1	129.309	9,009009009	1	5,487135172
A7E3D5	Proteasome subunit alpha type (Fragment)	26.551	4,856512141	0	10,87347269
K4JR81	Alpha-2-macroglobulin variant 12	61.673	4,856512141	3	10,87347269
Q3ZBG0	Proteasome subunit alpha type-7	27.852	8,411214953	2	126,5844053
F1MSZ6	Antithrombin-III OS=Bos taurus	52.407	11,02040816	9	126,5844053
P00735	Prothrombin	70.461	36,24733475	4	260,5843923
F1MFY6	Collectin-43 precursor	24.905	34,70437018	3	846,0876623
Q58D62	Fetuin-B	42.636	3,310613437	0	7,906161785
P06394	Keratin. type I cytoskeletal 10	54.815	6,204379562	2	6,61307168

Code access	Description*	MW [kDa]	Coverage	Unique peptides	Score
Q3SZJ0	Argininosuccinate lyase	52.71	4,744525547	0	6,086005092
G5E5C3	Proteasome subunit alpha type	27.267	10,57692308	1	4,239753962
F1MM86	Complement component C6	104.455	2,702702703	7	3,410375118
G3N0V2	Keratin 1	63.113	11,07954545	0	4,860876799
O02808	Von Willebrand factor (Fragment)	59.57	11,07954545	9	4,860876799
Q3SYT3	Complement C1r subcomponent precursor	44.528	5,113636364	10	7,11321342
Q2YDE4	Proteasome subunit alpha type-6	27.382	11,11111111	0	6,825133801
F1MTV7	Argininosuccinate lyase	52.682	11,11111111	1	6,825133801
B8YB77	Malic enzyme	63.332	5,812417437	4	7,035996199
G3X6N3	Serotransferrin	77.616	5,812417437	3	7,035996199
E1BFG1	Uncharacterized protein	50.005	4,756871036	0	11,78225541
F1MRZ6	Tenascin C	244.461	4,756871036	4	11,78225541
G5E5T5	Uncharacterized protein	42.442	5,113636364	1	7,11321342
Q3SYR5	Apolipoprotein C-IV	14.428	5,38573508	2	4,48806572
Q29RU4	Complement component C6	104.473	23,07692308	1	11,25674486
Q3T063	Nicotinate-nucleotide pyrophosphorylase	31.131	4,347826087	2	4,48806572
E1BL29	Bleomycin hydrolase	52.896	6,532663317	0	27,89325523
Q29443	Serotransferrin	77.703	20,15503876	2	3,950842738
F1MHB8	Nicotinate-nucleotide pyrophosphorylase	31.165	8,558558559	2	1,680921078
P41361	Antithrombin-III	52.314	12,5	1	110,3163168
Q71U44	Fibronectin (Fragment)	45.802	17,59656652	1	109,0141351
A0JN60	Tenascin C	190.961	20,47244094	2	5,039208889
E1BH94	Peptidoglycan recognition protein 2	59.793	23,95833333	0	7,306949139
F1MRZ5	Tenascin C	191.032	6,28742515	2	4,237655163
Q2KIV9	Complement C1q subcomponent subunit B	26.383	6,290672451	0	7,969627857
Q28194	Thrombospondin-1	25	1,493428913	2	4,015279174
Q9TUQ0	Anion exchange protein	95.582	6,982543641	10	2,994791746
Q148H5	Keratin. type II cytoskeletal	57.372	7,00525394	2	16,71615529
Q32KL2	Proteasome subunit beta type-5	28.591	7,00525394	6	16,71615529
P63258	Actin. cytoplasmic 2	41.766	4,22832981	2	4,367512226
B5B3R8	Alpha S1 casein	24.427	4,22832981	0	4,367512226
A0A1B0Z542	Heat shock 27 kDa protein 1	22.231	1,431127013	3	4,702090144
Q9XSW5	Anion exchange protein	104.308	5,735660848	0	9,489186764

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F1MNW4	Inter-alpha-trypsin inhibitor heavy chain H2	106.09	5,735660848	3	9,489186764
Q3MHF7	S-methyl-5'-thioadenosine phosphorylase	31.236	3,565062389	2	4,589993954
F1N3V0	Malic enzyme	63.746	3,584229391	1	4,589993954
F1MYN5	Fibulin-1	77.434	2,253855279	0	4,057821274
A5D7S8	Fibulin-1	77.478	2,253855279	3	4,057821274
A0A1C9EIX3	Heat shock protein family B member 1 variant 1	22.351	2,549575071	12	4,26041913
Q58DP7	Heat shock 27kDa protein 1	17.545	2,549575071	0	4,26041913
E9RHW1	Heat shock 27kDa protein 1	22.379	18,46153846	1	6,605324507
Q3T0X5	Proteasome subunit alpha type-1	29.567	18,46153846	0	6,605324507
P02662	Alpha-S1-casein	24.513	0,788643533	0	7,681156635
P60712	Actin. cytoplasmic 1	41.71	1,947419669	2	7,787001133
P31976	Ezrin	68.717	7,733333333	1	7,858686447
Q3ZCK9	Proteasome subunit alpha type-4	29.465	7,733333333	0	7,858686447
Q9TUQ1	Anion exchange protein	74.3	7,733333333	5	7,858686447
A0A1C9EIX6	Heat shock protein family B member 1 variant 2	22.389	1,431127013	3	4,702090144
A5D984	Pyruvate kinase	57.912	15,10791367	1	11,06255352
Q28085	Complement factor H	140.282	12,93103448	2	12,63318157
F1MC45	Complement factor H precursor	96.53	13,67521364	3	12,63318157
F1MRD0	Actin. cytoplasmic 1	41.825	10	2	12,63318157
Q9BGI2	Peroxiredoxin-4	30.722	12,76595744	0	12,63318157
Q3ZC87	Pyruvate kinase (Fragment)	61.389	27,06222865	1	453,5852154
E1BEL7	Heat shock protein beta-1	22.564	32,60869565	1	5,436547518
Q7M2T6	Band 3 anion transport protein (Fragments)	34.3	32,78688525	2	5,436547518
A5D7R6	ITIH2 protein	106.12	19,53125	1	5,436547518
G3X7S2	Heat shock protein beta-1	17.432	32,78688525	1	5,436547518
Q1JQB0	Collagen type VI alpha 2 chain	97.075	9,693877551	1	5,790735483
Q6T182	Sex hormone-binding globulin (Fragment)	40.069	6,181015453	0	6,662236333
Q29RQ1	Complement component C7	93.029	19,35483871	2	12,07949877
G5E5H7	Uncharacterized protein	19.898	16,02564103	0	5,436547518
K4JF16	Alpha-2-macroglobulin variant 23	101.296	16,25615764	1	21,39780414
Q2HJ86	Tubulin alpha-1D chain	50.251	3,434610304	3	4,737163544
P81947	Tubulin alpha-1B chain	50.12	2,599814299	5	2,994791746
A3KLR9	Superoxide dismutase [Cu-Zn]	26.16	2,094240838	0	2,994791746

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E1BFN5	Uncharacterized protein	78.333	24,12962557	1	674,8379145
Q1RML9	Platelet-activating factor acetylhydrolas	50.119	33,65442505	1	440,4326642
Q05B55	IGK protein	26.574	33,23299217	1	440,4326642
Q32KN8	Tubulin alpha-3 chain	49.894	32,69114991	2	410,7775776
Q6KDN5	Complement component 3 (Fragment)	7.414	27,06325932	2	327,4399574
F1MNV5	Kininogen-1	48.391	25,14124294	0	321,4439662
Q17QL7	KRT15 protein	48.969	25,14124294	0	321,4439662
K4JR88	Alpha-2-macroglobulin variant 22	77.345	26,54320988	2	318,3170657
B5B0D4	Major allergen beta-lactoglobulin	19.956	19,7564276	2	220,1763519
F1MH40	Uncharacterized protein	26.318	19,72972973	1	220,1763519
K4JDR8	Alpha-2-macroglobulin variant 5	45.046	59,62264151	2	232,1050236
A1L595	Keratin. type I cytoskeletal 17	48.682	8,790035587	2	115,8492713
A0A140T8C8	Kininogen-1	68.922	8,796296296	2	115,8492713
P01044	Kininogen-1	68.847	27,22371968	0	404,0782466
G3X6I0	Uncharacterized protein	202.852	26,1589404	3	873,9068512
F1N045	Complement component C7	92.929	14,44444444	2	65,70898986
Q2HJ49	Moesin	67.933	22,73476112	0	73,49145103
F2Z4C1	Tubulin alpha chain	50.104	22,73476112	2	73,49145103
P81644	Apolipoprotein A-II	11.195	48,55769231	3	119,6688337
A0A140T867	Keratin. type I cytoskeletal 17	48.712	15,57377049	2	100,6843615
Q32LP2	Radixin	68.525	13,41880342	3	62,5247829
F1MMP5	Inter-alpha-trypsin inhibitor heavy chain H1	101.174	30,3030303	0	52,56129336
A5PKC2	SHBG protein	43.288	32,05128205	2	52,56129336
P56652	Inter-alpha-trypsin inhibitor heavy chain H3	99.489	20,59308072	1	62,91113496
F6RP72	Tubulin alpha chain	49.797	13,60655738	2	100,6843615
Q28017	Platelet-activating factor acetylhydrolas	50.101	44,6685879	1	108,6491276
O46415	Ferritin light chain	19.975	25,67324955	0	303,3408718
M0QVY0	Uncharacterized protein	60.767	8,737864078	1	71,69110668
E1BD83	Proteasome subunit alpha type	29.184	19,0070922	1	31,68743467
O02717	Non-muscle myosin heavy chain (Fragment)	72.327	13,95749569	1	111,2448874
Q27991	Myosin-10	228.958	26,31578947	1	33,63287568
P01966	Hemoglobin subunit alpha	15.175	26,31578947	1	33,63287568
G5E604	Uncharacterized protein	11.051	30,0330033	3	76,51224542

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P02754	Beta-lactoglobulin	19.87	12,44239631	1	25,84321988
K4JBA2	Alpha-2-macroglobulin variant 9	43.75	24,29048414	2	646,0610726
A4FV94	KRT6A protein	60.783	6,357388316	0	20,59954548
B0JYP6	IGK protein	26.304	6,357388316	1	20,59954548
K4JBR5	Alpha-2-macroglobulin variant 1	115.118	4,975347378	0	20,59954548
Q0VCM5	Inter-alpha-trypsin inhibitor heavy chain H1	101.173	38,75	0	71,74341667
P01045	Kininogen-2	68.666	14,77477477	1	62,25517595
Q2TBX6	Proteasome subunit beta type-1	26.229	13,49593496	0	39,98661864
Q05443	Lumican	38.732	13,49593496	2	39,98661864
G3N0S9	Uncharacterized protein	22.321	27,84810127	1	25,54059851
P34955	Alpha-1-antiproteinase	46.075	27,84810127	1	25,54059851
P19035	Apolipoprotein C-III	10.685	27,84810127	1	25,54059851
F1MZ96	Uncharacterized protein	26.545	28,48484848	1	30,96007919
P33672	Proteasome subunit beta type-3	22.977	28,18627451	2	267,9672889
E1BB91	Collagen type VI alpha 3 chain	342.197	28,96725441	1	267,9672889
F2Z4K0	Tubulin alpha chain	49.928	24,31289641	1	267,9672889
P63103	14-3-3 protein zeta/delta	27.728	10,21582734	1	32,74262905
F1MJJ8	Radixin	68.541	10,30478955	4	32,74262905
G5E589	Proteasome subunit beta type	26.301	14,4469526	0	38,56606507
Q3ZCJ7	Tubulin alpha-1C chain	49.825	14,71264368	1	38,56606507
Q32LE5	Isoaspartyl peptidase/L-asparaginase	32.03	27,154047	2	227,8457786
K4JDS3	Alpha-2-macroglobulin variant 10	52.304	17,30418944	2	42,44953275
Q687I9	Purine nucleoside phosphorylase	32.046	36,24454148	1	32,37882257
F1MVJ8	Olfactomedin 4	57.76	25,43520309	0	505,1839079
F1MJH1	Gelsolin	80.653	32,38512035	1	770,8895055
G3X8C8	Uncharacterized protein	25.166	22,92993631	0	60,16956151
P62739	Actin, aortic smooth muscle	41.982	5,501618123	4	11,86303473
Q28921	Alpha 1-antichymotrypsin (Fragment)	28.553	13,30798479	3	32,08415091
Q28007	Dihydropyrimidine dehydrogenase [NADP(+)]	111.625	13,30798479	2	32,08415091
Q3SX14	Gelsolin	80.681	31,64556962	1	14,37422311
Q9XTA3	Myocilin	54.853	11,03603604	1	33,17564273
A2I7M9	Serpin A3-2	46.208	27,27272727	1	148,6272044
B0JYK6	Alpha-1.4 glucan phosphorylase	97.227	20,38834951	2	65,30266976

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Q1JPD0	Complement component 8. alpha polypeptide	32.621	25,41666667	1	22,7700392
P80012	von Willebrand factor (Fragment)	102.531	10,93394077	1	23,2725662
Q862P9	Similar to beta actin (Fragment)	20.968	12,14128035	1	87,20624363
F1MKC4	Actin. gamma-enteric smooth muscle	41.898	12,19512195	0	87,20624363
Q2KIH5	Complement component 8. alpha polypeptide	66.292	10,54347826	3	51,87173843
G3MXU3	Collagen type VI alpha 2 chain	24.991	28,76106195	2	551,6399485
G3N126	Collagen type VI alpha 3 chain	134.797	8,333333333	0	14,14523625
A0A140T871	Glutamate dehydrogenase 1. mitochondrial	61.592	17,81376518	1	26,02097392
Q58DU5	Proteasome subunit alpha type-3	28.387	12,40875912	4	23,08411241
F1MJ28	Alpha-1.4 glucan phosphorylase	97.218	7,755775578	6	24,12152457
Q3T186	Ribose-5-phosphate isomerase	28.735	48,59813084	3	65,37755024
Q8SPJ1	Junction plakoglobin	81.769	5,176767677	0	9,596704483
A6QQC9	OLFM4 protein (Fragment)	49.706	12,43654822	2	11,27226317
Q28908	Mucin (Fragment)	54.83	9,009009009	2	5,487135172
F1MX87	Complement C8 alpha chain	66.234	9,009009009	1	5,487135172
H9KUV2	S-methyl-5'-thioadenosine phosphorylase	32.883	4,856512141	0	10,87347269
G3N1U4	Serpin A3-3	46.127	4,856512141	1	10,87347269
F1MNF8	Tubulin alpha chain	49.87	8,411214953	1	126,5844053
A3FJ56	Kappa casein (Fragment)	17.836	11,02040816	1	126,5844053
Q705V4	Kappa-casein (Fragment)	17.693	36,24733475	0	260,5843923
Q0VCX2	Endoplasmic reticulum chaperone BiP	72.356	34,70437018	3	846,0876623
A5D7L1	C-type lectin domain containing 11A	35.593	3,310613437	0	7,906161785
Q3T052	Inter-alpha-trypsin inhibitor heavy chain H4	101.449	6,204379562	2	6,61307168
F1N549	Dihydropyrimidine dehydrogenas	111.766	4,744525547	4	6,086005092
G5E534	Ribose-5-phosphate isomerase	32.82	10,57692308	3	4,239753962
Q3ZC07	Actin. alpha cardiac muscle	41.992	2,702702703	2	3,410375118
Q2KJF1	Alpha-1B-glycoprotein	53.52	11,07954545	3	4,860876799
Q5EA67	Inter-alpha (Globulin) inhibitor H4 (Plasma Kallikrein-sensitive glycoprotein)	101.446	11,07954545	2	4,860876799
Q3B7M9	Glycogen phosphorylase	96.279	5,113636364	1	7,11321342
F1N1I6	Gelsolin	85.634	11,11111111	3	6,825133801
O46375	Transthyretin	15.717	11,11111111	5	6,825133801
F1N0I3	Coagulation factor V	222.078	5,812417437	3	7,035996199

Code access	Description*	MW [kDa]	Coverage	Unique peptides	Score
P02081	Hemoglobin fetal subunit beta	15.849	5,812417437	2	7,035996199
P02668	Kappa-casein	21.256	4,756871036	3	11,78225541
B8YB76	Homogentisat	49.965	4,756871036	2	11,78225541
Q9TTE1	Serpin A3-1	46.208	5,113636364	3	7,11321342
P81948	Tubulin alpha-4A chain	49.892	5,38573508	2	4,48806572
Q3SX06	Myocilin	54.854	23,07692308	3	11,25674486
F1MU24	Alpha-1.4 glucan phosphorylase	76.125	4,347826087	3	4,48806572
P68138	Actin. alpha skeletal muscle	42.024	6,532663317	2	27,89325523
F1MXQ3	FAM20C. golgi associated secretory pathway kinase	61.848	20,15503876	1	3,950842738
A8YXZ2	C8G protein	25.272	8,558558559	2	1,680921078
A2I7N1	Serpin A3-5	46.368	12,5	0	110,3163168
Q3ZBS7	Vitronectin	53.541	17,59656652	1	109,0141351
D4QBB4	Globin A1	15.944	20,47244094	1	5,039208889
A5D7Q2	Uncharacterized protein	51.638	23,95833333	1	7,306949139
M0QVZ6	Keratin. type II cytoskeletal 5	60.629	6,28742515	1	4,237655163
Q2KIH3	HGD protein (Fragment)	44.278	6,290672451	1	7,969627857
Q5E9B5	Actin. gamma-enteric smooth muscle	41.85	1,493428913	1	4,015279174
Q3ZEJ6	Serpin A3-3	46.297	6,982543641	1	2,994791746
Q58DT9	Alpha 2 actin	45.192	7,00525394	1	16,71615529
G3N1Y3	Uncharacterized protein	12.966	7,00525394	1	16,71615529
G3MX98	Keratin 9	54.621	4,22832981	1	4,367512226
Q9MYV8	Haptoglobin (Fragment)	11.232	4,22832981	1	4,367512226
G3N3E4	Collagen type VI alpha 3 chain	185.721	1,431127013	1	4,702090144
Q32L76	Serum amyloid A-4 protein	14.678	5,735660848	1	9,489186764
G8JKX4	Actin. aortic smooth muscle	45.403	5,735660848	1	9,489186764
A0A0M4FJ17	Kappa-casein (Fragment)	16.015	3,565062389	1	4,589993954
P55859	Purine nucleoside phosphorylase	32.016	3,584229391	1	4,589993954
F1N614	78 kDa glucose-regulated protein precursor	66.224	2,253855279	1	4,057821274
A5D7M6	KRT5 protein	62.644	2,253855279	1	4,057821274
Q27983	Alpha1-antichymotrypsin isoform pHHK11 (Fragment)	22.697	2,549575071	1	4,26041913
A6QPD4	LOC790886 protein	45.399	2,549575071	1	4,26041913
F1MMD7	Inter-alpha-trypsin inhibitor heavy chain H4	101.463	18,46153846	1	6,605324507
A4IFM8	Actin. alpha 1. skeletal muscle	41.996	18,46153846	1	6,605324507

Code access	Description*	MW [kDa]	Coverage	Unique peptides	Score
A2I7N0	Serpin A3-4	46.282	0,788643533	1	7,681156635
F1MLF8	Homogentisat	49.811	1,947419669	1	7,787001133
P81265	Polymeric immunoglobulin receptor	82.383	7,733333333	1	7,858686447
A2I7N2	Serpin A3-6	46.361	7,733333333	1	7,858686447
K4JB97	Alpha-2-macroglobulin variant 4	42.306	7,733333333	1	7,858686447
A0A140T8A9	Kappa-casein	21.224	1,431127013	1	4,702090144
Q28107	Coagulation factor V	248.828	15,10791367	1	11,06255352
Q0P5J4	Keratin. type I cytoskeletal 25	49.282	10	1	12,63318157
A5D7J0	ACTA2 protein	42.01	10	1	12,63318157
P00366	Glutamate dehydrogenas	61.473	10	1	12,63318157
Q3ZC00	Lymphocyte cytosolic protein 1 (L-plastin)	70.01	10	1	12,63318157
Q148H6	Keratin. type I cytoskeletal 28	50.744	27,06222865	1	453,5852154
G8JKW7	Uncharacterized protein	46.315	32,60869565	1	5,436547518
Q9MZ31	Fibronectin variable region (Fragment)	20.365	32,78688525	1	5,436547518
Q2HJB8	Tubulin alpha-8 chain	50.022	19,53125	1	5,436547518
F1MYX5	Lymphocyte cytosolic protein 1	70.068	32,78688525	1	5,436547518
A6QNW3	PIGR protein	82.455	9,693877551	1	5,790735483
Q5XQN5	Keratin. type II cytoskeletal 5	62.898	6,181015453	1	6,662236333
P79334	Glycogen phosphorylase	97.232	19,35483871	1	12,07949877
Q3MHN2	Complement component C9	61.958	16,02564103	1	5,436547518

*Classification given by author.