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

Additional file 2. Summary of extracellular vesicle proteins of *Theileria* spp. isolated from serum of control animals present in Fraction F3.

Code access	Description*	MW [kDa]	Coverage	Unique Peptides	Score
E1BNR0	Apolipoprotein	515.438	23.40705	94	495.0726119
Q2UVX4	Complement C3	187.135	31.66767	52	345.5245875
A0A0F6QNP7	Complement component 3	187.064	31.24623	51	343.8403611
G3X7A5	Complement C3	187.028	30.70439	51	330.6581188
B8Y9S9	Fibronectin 1	262.263	15.37494	24	110.1794619
G5E5A9	Fibronectin	271.952	14.81033	24	110.1794619
P07589	Fibronectin	271.983	14.81033	24	110.1794619
B8Y9T0	Fibronectin 1	248.974	16.18165	24	110.1794619
P15497	Apolipoprotein A-I	30.258	47.16981	15	117.2911120
Q0VCM4	Glycogen phosphorylase	97.394	1.527614	1	0
F1MI18	Uncharacterized protein	165.654	19.68876	26	176.92844569
F1MJK3	Uncharacterized protein	165.506	19.66216	26	176.9284456
A0A140T897	Serum albumin	69.278	28.33607	16	87.0102145
P02769	Serum albumin	69.248	28.33607	16	87.0102145
P23805	Conglutinin	37.971	26.68463	14	380.1768126
P80457	Xanthine dehydrogenase/oxidase	146.696	0.750750	1	2.965109582
F1MUT3	Xanthine dehydrogenase/oxidase	146.669	0.750750	16	2.965109586
B0JYQ0	ALB protein	69.248	26.19439	15	80.409170196
F1MVP0	ADAM metallopeptidase with thrombospondin type 1	151.374	19.67213	15	120.12379646
F1N0R5	von Willebrand factor	307.475	3.596866	8	26.1033883094
F1MAV0	Fibrinogen beta chain	56.405	20.40404	8	51.431623342
E1BH06	Uncharacterized protein	192.644	15.62320	10	86.6288429
F5XVA9	von Willebrand factor	307.745	3.594306	8	26.10338830

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P02676	Fibrinogen beta chain	53.306	20.08547	7	25.59032678
F1MY85	Complement C5a anaphylatoxin	188.927	2.444841	4	8.5798010826
V6F869	Apolipoprotein A-I-like	23.629	29.80769	7	44.303140
A0A0F6QMJ3	Complement component 5	188.691	2.444841	4	8.5798010
F1N3Q7	Apolipoprotein A-IV	42.963	30.52631	13	71.981267
Q32PJ2	Apolipoprotein A-IV	42.991	30.52631	13	71.981267
G3N0I4	Cytosol aminopeptidas	54.047	1.606425	1	1.9673806
P00727	Cytosol aminopeptidas	56.254	1.967380	1	1.9673802
Q693V9	Complement component 3d	34.422	21.45214	1	87.169953
Q3MHL4	Adenosylhomocysteinase 3	47.607	21.40992	7	137.36276
P00432	Catalase	59.878	2.466793	1	0
Q7SIH1	Alpha-2-macroglobulin	167.47	31.32450	10	668.51652
Q95KV5	Fibronectin	38.32	38.04034	9	46.092858
F1MQ37	Myosin heavy chain 9	226.962	19.67213	13	134.312602
F1MD73	Uncharacterized protein	189.978	12.50713	10	35.2686412
A5PJE3	Fibrinogen alpha chain	66.957	10.08130	6	18.8123501
P02672	Fibrinogen alpha chain	66.971	10.08130	6	18.8123501
A5D9E9	Complement C1r subcomponent precursor	80.161	5.673758	1	6.06269646
Q03247	Apolipoprotein	35.958	21.51898	6	50.4440462
Q2TBU0	Haptoglobin	44.831	23.19201	8	48.0335323
E1BI98	Collagen type VI alpha 1 chain	108.603	21.867321	6	124.459278
G5E513	Uncharacterized protein	49.939	20	6	22.3382167
A6QPX7	FGB protein (Fragment)	37.907	9.456521	5	35.100500
P01030	Complement C4	101.817	17.06522	9	43.73454607
E1B805	Uncharacterized protein	187.149	23.192019	8	48.0335323
G3X6K8	Haptoglobin	44.845	25.69327	8	34.74552003
A0A140T881	Apolipoprotein	36.017	21.518987	9	50.4440462
G3N3D4	Potassium channel tetramerization domain containing 12	35.701	8.5427135	3	41.012642
A7YWR0	Apolipoprotein	36.051	21.51898	3	21.51898
E1BFN6	Dihydropyrimidinase	56.267	4.207119	5	4.20711974
P81187	Complement factor B	85.312	1.7241379	1	4.834334
P17697	Clusterin	51.081	9.1116173	3	10.1167018
F1N076	Ceruloplasmin	123.731	4.792626	4	12.1857370

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R9QSM8	Alpha-2-macroglobulin	133.302	32.637729	10	531.153768
F1N514	CD5 antigen-like precursor	50.305	20.75055	7	16.0244067
Q3SZZ9	FGG protein	49.136	18.16091	8	38.8492791
Q0VCX1	Complement C1s subcomponent	76.56	6.386066	12	12.067846
F1MJ12	Complement C1s subcomponent	77.332	6.330935	12	23.25133181
F1MGU7	Fibrinogen gamma-B chain	50.2	17.83295	8	38.8492790
Q95KV4	Fibronectin (Fragment)	35.464	5.184630	4	5.97265890
Q3Y5Z3	Adiponectin	26.117	20.74950	8	29.8565749
Q0ZCB4	Apolipoprotein	27.057	7.9755563	3	0.98859376
A2VDY5	Hydroxysteroid (17-beta)	28.378	8.1481481	2	4.8307673
Q9MYP6	17-beta-hydroxysteroid dehydrogenas	28.404	24.27707	3	9.39274930
A6QNW7	CD5L protein	50.179	16.837208	4	8.77830857
Q0P5J7	Keratin. type I cuticular Ha5	49.905	1.538461	2	2.12892317
B7FEK7	43kDa collectin	33.594	8.411214	4	129.4997527
E1BPW6	Keratin. type I cuticular Ha5	49.739	1.54867	2	2.12892317
Q28178	Thrombospondin-1	129.451	10.2564102	11	66.87595036
A7E3W2	Galectin-3-binding protein	62.087	8.6486486	5	26.75749302
F1MKG2	Collagen type VI alpha 2 chain	109.253	2.0447994	2	5.74746538
B8Y898	Malic enzyme	63.853	12.8640776	3	28.8594936
P02663	Alpha-S2-casein	26.002	8.5585585	2	6.87464993
F1MC11	Keratin. type I cytoskeletal 14	51.879	15.72327	6	28.12601042
A6QNZ7	Keratin 10	54.816	8.55513	4	31.8585664
P12799	Fibrinogen gamma-B chain	50.212	1.441441	7	34.8575392
Q3SYR8	Immunoglobulin J chain	17.846	22.92993	6	69.2885639
F1N3A1	Thrombospondin-1	129.309	11.282051	1	68.986869
A7E3D5	Proteasome subunit alpha type (Fragment)	26.551	4.6413502	1	1.7184752
K4JR81	Alpha-2-macroglobulin variant 12	61.673	21.723518	11	186.688745
Q3ZBG0	Proteasome subunit alpha type-7	27.852	4.4354838	1	1.88745493
F1MSZ6	Antithrombin-III OS=Bos taurus	52.407	10.96774	4	15.8486339
P00735	Prothrombin	70.461	3.27510	1	7.94946922
F1MFY6	Collectin-43 precursor	24.905	11.02040	4	129.993746
Q58D62	Fetuin-B	42.636	8.485856	3	41.0936353
P06394	Keratin. type I cytoskeletal 10	54.815	8.555133	4	31.9936534

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Q3SZJ0	Argininosuccinate lyase	52.71	10.847453	8	120.948452
G5E5C3	Proteasome subunit alpha type	27.267	10	4	7.99463420
F1MM86	Complement component C6	104.455	5.955242	3	0.99374294
G3N0V2	Keratin 1	63.113	9.075907	5	43.8835422
O02808	Von Willebrand factor (Fragment)	59.57	12.99825	5	130.999273
Q3SYT3	Complement C1r subcomponent precursor	44.528	12.992654	4	15.88864435
Q2YDE4	Proteasome subunit alpha type-6	27.382	20.993752	3	12.89826425
F1MTV7	Argininosuccinate lyase	52.682	2.1515015	4	238.998765
B8YB77	Malic enzyme	63.332	5.9972543	5	10.993764
G3X6N3	Serotransferrin	77.616	7.9864311	5	9.9864321
E1BFG1	Uncharacterized protein	50.005	15.23342	2	2.12892319
F1MRZ6	Tenascin C	244.461	4.9987354	3	0.987354335
G5E5T5	Uncharacterized protein	42.442	4.997353	1	0.09375343
Q3SYR5	Apolipoprotein C-IV	14.428	8.0937635	1	0.98365343
Q29RU4	Complement component C6	104.473	12.0937354	5	129.098765
Q3T063	Nicotinate-nucleotide pyrophosphorylase	31.131	3.983754	4	6.09487465
E1BL29	Bleomycin hydrolase	52.896	9.98476453	4	308.948743
Q29443	Serotransferrin	77.703	6.9876543	5	21.947465443
F1MHB8	Nicotinate-nucleotide pyrophosphorylase	31.165	10.0948476	1	0.98765438
P41361	Antithrombin-III	52.314	12.094765	3	11.0946544
Q71U44	Fibronectin (Fragment)	45.802	15.96117	4	16.0987654
A0JN60	Tenascin C	190.961	3.180985	2	1.98476544
E1BH94	Peptidoglycan recognition protein 2	59.793	12.0958765	3	11.5987654
F1MRZ5	Tenascin C	191.032	3.0958765	1	7.59876543
Q2KIV9	Complement C1q subcomponent subunit B	26.383	7.0948765	1	7.09586532
Q28194	Thrombospondin-1	25	1.98576543	2	5.09765432
Q9TUQ0	Anion exchange protein	95.582	3.987654	3	11.87643221
Q148H5	Keratin. type II cytoskeletal	57.372	1.714285	1	16.02440679
Q32KL2	Proteasome subunit beta type-5	28.591	15.887654	2	5.875421187
P63258	Actin. cytoplasmic 2	41.766	10.987653	3	9.876543215
B5B3R8	Alpha S1 casein	24.427	9.9876543	3	13.87654321
A0A1B0Z542	Heat shock 27 kDa protein 1	22.231	12.987654	1	0.09876543
Q9XSW5	Anion exchange protein	104.308	2.8765432	4	15.87643221

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F1MNW4	Inter-alpha-trypsin inhibitor heavy chain H2	106.09	4.75687103	3	6.98654432
Q3MHF7	S-methyl-5'-thioadenosine phosphorylase	31.236	19.986432	3	7.9876543
F1N3V0	Malic enzyme	63.746	11.986432	3	12.8765432
F1MYN5	Fibulin-1	77.434	8.087654	3	8.98765454
A5D7S8	Fibulin-1	77.478	2.987654	2	3.9876544
A0A1C9EIX3	Heat shock protein family B member 1 variant 1	22.351	3.0987654	2	8.98765438
Q58DP7	Heat shock 27kDa protein 1	17.545	10.987654	3	9.98765434
E9RHW1	Heat shock 27kDa protein 1	22.379	9.9876543	2	7.9876543
Q3T0X5	Proteasome subunit alpha type-1	29.567	43.28358	4	238.9876543
P02662	Alpha-S1-casein	24.513	11.682242	2	4.09876548
P60712	Actin. cytoplasmic 1	41.71	7.7333333	2	7.9876924
P31976	Ezrin	68.717	6.9876543	2	4.0898243
Q3ZCK9	Proteasome subunit alpha type-4	29.465	19.9876543	4	27.9289973
Q9TUQ1	Anion exchange protein	74.3	5.987654	4	9.763430
A0A1C9EIX6	Heat shock protein family B member 1 variant 2	22.389	19.998533	3	5.9876543
A5D984	Pyruvate kinase	57.912	10.998745	3	120.87654
Q28085	Complement factor H	140.282	4.9988543	3	3.87654365
F1MC45	Complement factor H precursor	96.53	9.993425	3	10.987654
F1MRD0	Actin. cytoplasmic 1	41.825	10.096543	2	13.9937432
Q9BGI2	Peroxiredoxin-4	30.722	18.1983738	4	120.09376354
Q3ZC87	Pyruvate kinase (Fragment)	61.389	1.0987654	2	0.987650876
E1BEL7	Heat shock protein beta-1	22.564	2.021975	2	8.093763543
Q7M2T6	Band 3 anion transport protein (Fragments)	34.3	10.983753	4	186.09837365
A5D7R6	ITIH2 protein	106.12	2.0987643	2	1.987354332
G3X7S2	Heat shock protein beta-1	17.432	4.0876524	3	4.08754221
Q1JQB0	Collagen type VI alpha 2 chain	97.075	2.0976543	1	0.98765432
Q6T182	Sex hormone-binding globulin (Fragment)	40.069	10.087654	2	7.0987643
Q29RQ1	Complement component C7	93.029	2.9876543	3	12.9876543
G5E5H7	Uncharacterized protein	19.898	23.987654	1	4.9876543
K4JF16	Alpha-2-macroglobulin variant 23	101.296	5.09871457	1	19.8632145
Q2HJ86	Tubulin alpha-1D chain	50.251	19.9876543	1	3.986439876
P81947	Tubulin alpha-1B chain	50.12	3.9876543	1	8.09876543
A3KLR9	Superoxide dismutase [Cu-Zn]	26.16	12.9876543	1	10.87654987

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E1BFN5	Uncharacterized protein	78.333	10.0987654	2	5.0882533
Q1RML9	Platelet-activating factor acetylhydrolas	50.119	12.0937343	1	0.98764322
Q05B55	IGK protein	26.574	10.0987654	1	11.0987654
Q32KN8	Tubulin alpha-3 chain	49.894	9.0987654	2	6.0947542
Q6KDN5	Complement component 3 (Fragment)	7.414	47.094876	2	0.09464367
F1MNV5	Kininogen-1	48.391	5.984454	3	28.09476543
Q17QL7	KRT15 protein	48.969	12.0987654	3	110.1794619
K4JR88	Alpha-2-macroglobulin variant 22	77.345	4.0984326	1	6.2589853
B5B0D4	Major allergen beta-lactoglobulin	19.956	2.986421	2	0.986422467
F1MH40	Uncharacterized protein	26.318	2.76542345	1	1.09876411
K4JDR8	Alpha-2-macroglobulin variant 5	45.046	5.8765431	1	0.63212345
A1L595	Keratin. type I cytoskeletal 17	48.682	7.7654234	1	0.95327888
A0A140T8C8	Kininogen-1	68.922	5.9876543	1	3.9876543
P01044	Kininogen-1	68.847	5.09876	1	0.14987654
G3X6I0	Uncharacterized protein	202.852	1.9876543	1	0.260093635
F1N045	Complement component C7	92.929	0.677343	2	12.8753321
Q2HJ49	Moesin	67.933	8.0846354	1	7.05987364
F2Z4C1	Tubulin alpha chain	50.104	7.09483765	1	6.50948736
P81644	Apolipoprotein A-II	11.195	2	1	0.16497433
A0A140T867	Keratin. type I cytoskeletal 17	48.712	3.594306	1	6.0924317
Q32LP2	Radixin	68.525	1.059876	1	2.0598473
F1MMP5	Inter-alpha-trypsin inhibitor heavy chain H1	101.174	3.098541	1	2.93753365
A5PKC2	SHBG protein	43.288	11.0847354	2	31.0475233
P56652	Inter-alpha-trypsin inhibitor heavy chain H3	99.489	1.0847354	2	5.05948736
F6RP72	Tubulin alpha chain	49.797	2.054654	1	2.059847365
Q28017	Platelet-activating factor acetylhydrolas	50.101	3.0948376	1	0.0594873
O46415	Ferritin light chain	19.975	32.059487	0	0.0598476
M0QVY0	Uncharacterized protein	60.767	2.0984634	1	0.02862225
E1BD83	Proteasome subunit alpha type	29.184	15.038634	1	11.34855783
O02717	Non-muscle myosin heavy chain (Fragment)	72.327	6.0947432	1	7.98362433
Q27991	Myosin-10	228.958	0.0598476	1	4.958735433
P01966	Hemoglobin subunit alpha	15.175	3.5167384	1	8.0594764
G5E604	Uncharacterized protein	11.051	7.04752456	1	69.0584635

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P02754	Beta-lactoglobulin	19.87	5.4	1	0.0918625
K4JBA2	Alpha-2-macroglobulin variant 9	43.75	16.59574468	1	134.4589926
A4FV94	KRT6A protein	60.783	2.272727273	1	2.706366062
B0JYP6	IGK protein	26.304	4.166666667	1	14.54332852
K4JBR5	Alpha-2-macroglobulin variant 1	115.118	2.00729927	2	1.662858129
Q0VCM5	Inter-alpha-trypsin inhibitor heavy chain H1	101.173	20	1	4.905816555
P01045	Kininogen-2	68.666	12.93103448	1	17.31717205
Q2TBX6	Proteasome subunit beta type-1	26.229	2.466793169	1	0
Q05443	Lumican	38.732	17.09401709	1	138.1611656
G3N0S9	Uncharacterized protein	22.321	1.642335766	1	2.366074562
P34955	Alpha-1-antiproteinase	46.075	3.284671533	1	2.431547165
P19035	Apolipoprotein C-III	10.685	1.527614571	1	0
F1MZ96	Uncharacterized protein	26.545	1.62601626	1	4.03635323
P33672	Proteasome subunit beta type-3	22.977	4.530744337	1	7.071590424
E1BB91	Collagen type VI alpha 3 chain	342.197	1.724137931	1	4.834334612
F2Z4K0	Tubulin alpha chain	49.928	3.643724696	1	1.762604713
P63103	14-3-3 protein zeta/delta	27.728	8.196721311	1	3.0535779
F1MJJ8	Radixin	68.541	3.542234332	1	2.346296549
G5E589	Proteasome subunit beta type	26.301	4.013377926	1	3.323835135
Q3ZCJ7	Tubulin alpha-1C chain	49.825	1.408450704	1	2.039117575
Q32LE5	Isoaspartyl peptidase/L-asparaginase	32.03	4.013377926	1	3.323835135
K4JDS3	Alpha-2-macroglobulin variant 10	52.304	1.269035533	1	2.039117575
Q687I9	Purine nucleoside phosphorylase	32.046	4.191616766	1	2.190802097
F1MVJ8	Olfactomedin 4	57.76	4.191616766	1	2.190802097
F1MJH1	Gelsolin	80.653	3.539823009	1	0
G3X8C8	Uncharacterized protein	25.166	3.921568627	1	0
P62739	Actin. aortic smooth muscle	41.982	0.750750751	1	2.965109587
Q28921	Alpha 1-antichymotrypsin (Fragment)	28.553	0.094746435	1	3.094763543
Q28007	Dihydropyrimidine dehydrogenase [NADP(+)]	111.625	18.8034188	1	14.98669171
Q3SX14	Gelsolin	80.681	15.17241379	1	14.98669171
Q9XTA3	Myocilin	54.853	6.986899563	1	4.565480947
A2I7M9	Serpin A3-2	46.208	9.142857143	1	4.565480947
B0JYK6	Alpha-1.4 glucan phosphorylase	97.227	5.97826087	1	2.060161829

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Q1JPD0	Complement component 8. alpha polypeptide	32.621	5.97826087	1	2.060161829
P80012	von Willebrand factor (Fragment)	102.531	3.375527426	1	1.675248504
Q862P9	Similar to beta actin (Fragment)	20.968	4.854368932	1	16.78249788
F1MKC4	Actin. gamma-enteric smooth muscle	41.898	4.862236629	1	16.78249788
Q2KIH5	Complement component 8. alpha polypeptide	66.292	2.330508475	1	1.985862494
G3MXU3	Collagen type VI alpha 2 chain	24.991	2.330508475	1	1.985862494
G3N126	Collagen type VI alpha 3 chain	134.797	31.25	1	0
A0A140T871	Glutamate dehydrogenase 1. mitochondrial	61.592	6.530612245	1	0
Q58DU5	Proteasome subunit alpha type-3	28.387	3.703703704	1	1.745364189
F1MJ28	Alpha-1.4 glucan phosphorylase	97.218	3.703703704	1	1.745364189
Q3T186	Ribose-5-phosphate isomerase	28.735	0.948991696	1	1.911735296
Q8SPJ1	Junction plakoglobin	81.769	0.948991696	1	1.911735296
A6QQC9	OLFM4 protein (Fragment)	49.706	1.146788991	1	0
Q28908	Mucin (Fragment)	54.83	1.606425703	1	1.967380643
F1MX87	Complement C8 alpha chain	66.234	1.541425819	1	1.967380643
H9KUV2	S-methyl-5'-thioadenosine phosphorylase	32.883	4.435483871	1	1.710465074
G3N1U4	Serpin A3-3	46.127	40	1	1.683246493
F1MNF8	Tubulin alpha chain	49.87	4.641350211	1	1.710465074
A3FJ56	Kappa casein (Fragment)	17.836	7.185628743	1	1.683246493
Q705V4	Kappa-casein (Fragment)	17.693	8.108108108	1	1.683246493
Q0VCX2	Endoplasmic reticulum chaperone BiP	72.356	4.214559387	1	1.710465074
A5D7L1	C-type lectin domain containing 11A	35.593	7.441860465	1	11.34865713
Q3T052	Inter-alpha-trypsin inhibitor heavy chain H4	101.449	7.441860465	1	11.34865713
F1N549	Dihydropyrimidine dehydrogenas	111.766	4.366812227	1	2.962756634
G5E534	Ribose-5-phosphate isomerase	32.82	39.36781609	1	177.1562679
Q3ZC07	Actin. alpha cardiac muscle	41.992	7.00525394	1	35.08573425
Q2KJF1	Alpha-1B-glycoprotein	53.52	7.00525394	1	35.08573425
Q5EA67	Inter-alpha (Globulin) inhibitor H4 (Plasma Kallikrein-sensitive glycoprotein)	101.446	22.5	1	2.310394764
Q3B7M9	Glycogen phosphorylase	96.279	1.181102362	1	2.475919485
F1N1I6	Gelsolin	85.634	10.5	1	2.576021194
O46375	Transthyretin	15.717	15.10791367	1	28.31982374
F1N0I3	Coagulation factor V	222.078	32.69012486	1	412.5393873

Code access	Description*	MW [kDa]	Coverage	Unique Peptides	Score
P02081	Hemoglobin fetal subunit beta	15.849	3.140096618	1	2.346296549
P02668	Kappa-casein	21.256	3.140096618	1	2.346296549
B8YB76	Homogentisat	49.965	1.449275362	1	1.972269893
Q9TTE1	Serpin A3-1	46.208	2.064220183	1	1.972269893
P81948	Tubulin alpha-4A chain	49.892	4.010025063	1	4.602294922
Q3SX06	Myocilin	54.854	3.925233645	1	21.29518712
F1MU24	Alpha-1.4 glucan phosphorylase	76.125	5.594405594	1	2.830938339
P68138	Actin. alpha skeletal muscle	42.024	4.929577465	1	1.777470112
F1MXQ3	FAM20C. golgi associated secretory pathway kinase	61.848	4.506437768	1	24.21742165
A8YXZ2	C8G protein	25.272	1.923076923	1	1.738595843
A2I7N1	Serpin A3-5	46.368	5.66893424	1	19.4173373
Q3ZBS7	Vitronectin	53.541	5.66893424	1	19.4173373
D4QBB4	Globin A1	15.944	5.353319058	1	19.4173373
A5D7Q2	Uncharacterized protein	51.638	7.00525394	1	35.08573425
M0QVZ6	Keratin. type II cytoskeletal 5	60.629	12.76595745	1	6.226527929
Q2KIH3	HGD protein (Fragment)	44.278	12.4137931	1	6.226527929
Q5E9B5	Actin. gamma-enteric smooth muscle	41.85	12.67605634	1	6.226527929
Q3ZEJ6	Serpin A3-3	46.297	13.63636364	1	6.226527929
Q58DT9	Alpha 2 actin	45.192	11.25	1	6.226527929
G3N1Y3	Uncharacterized protein	12.966	9.473684211	1	6.226527929
G3MX98	Keratin 9	54.621	12.85714286	1	6.226527929
Q9MYV8	Haptoglobin (Fragment)	11.232	11.32075472	1	6.226527929
G3N3E4	Collagen type VI alpha 3 chain	185.721	12.94964029	1	6.226527929
Q32L76	Serum amyloid A-4 protein	14.678	9.473684211	1	6.226527929
G8JKX4	Actin. aortic smooth muscle	45.403	13.23529412	1	6.226527929
A0A0M4FJ17	Kappa-casein (Fragment)	16.015	12.94964029	1	6.226527929
P55859	Purine nucleoside phosphorylase	32.016	1.201602136	1	2.475919485
F1N614	78 kDa glucose-regulated protein precursor	66.224	2.917771883	1	5.472857237
A5D7M6	KRT5 protein	62.644	2.925531915	1	5.472857237
Q27983	Alpha1-antichymotrypsin isoform pHHK11 (Fragment)	22.697	1.304347826	1	2.505360842
A6QPD4	LOC790886 protein	45.399	1.271186441	1	2.505360842
F1MMD7	Inter-alpha-trypsin inhibitor heavy chain H4	101.463	2.535211268	1	2.505360842
A4IFM8	Actin. alpha 1. skeletal muscle	41.996	1.271186441	1	2.505360842

Code access	Description*	MW [kDa]	Coverage	Unique Peptides	Score
A2I7N0	Serpin A3-4	46.282	5.102040816	1	0
F1MLF8	Homogentisat	49.811	1.199563795	1	2.096652985
P81265	Polymeric immunoglobulin receptor	82.383	8.064516129	1	6.581345797
A2I7N2	Serpin A3-6	46.361	0.858369099	1	0
K4JB97	Alpha-2-macroglobulin variant 4	42.306	0.858369099	1	0
A0A140T8A9	Kappa-casein	21.224	1.453957997	1	1.972269893
Q28107	Coagulation factor V	248.828	1.449275362	1	1.972269893
Q0P5J4	Keratin. type I cytoskeletal 25	49.282	5.617977528	1	26.10827029
A5D7J0	ACTA2 protein	42.01	4.929577465	1	1.777470112
P00366	Glutamate dehydrogenas	61.473	1.717305152	1	2.41721487
Q3ZC00	Lymphocyte cytosolic protein 1 (L-plastin)	70.01	2.41187384	1	2.41721487
Q148H6	Keratin. type I cytoskeletal 28	50.744	4.87012987	1	29.03050482
G8JKW7	Uncharacterized protein	46.315	2.917771883	1	5.472857237
Q9MZ31	Fibronectin variable region (Fragment)	20.365	1.923076923	1	1.738595843
Q2HJB8	Tubulin alpha-8 chain	50.022	1.19047619	1	2.475919485
F1MYX5	Lymphocyte cytosolic protein 1	70.068	3.867403315	1	24.21742165
A6QNW3	PIGR protein	82.455	39.408867	1	187.7336295
Q5XQN5	Keratin. type II cytoskeletal 5	62.898	2.709359606	1	5.472857237
P79334	Glycogen phosphorylase	97.232	2.925531915	1	5.472857237
Q3MHN2	Complement component C9	61.958	2.917771883	1	5.472857237

*Classification given by author.