

Summary Protein Identifications

#	Accession	IEF-PAGE	Blue Native	Tricine-Urea	16-BAC	No. of Methods ^a	Protein
1	gi 27806769	+	+	+	+	****	aconitase 2, mitochondrial [Bos taurus]
2	gi 27807143	+	+	+	+	****	ubiquinol-cytochrome c reductase core protein II [Bos taurus]
3	gi 27807237	+	+	+	+	****	ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit [Bos taurus]
4	gi 27807355	+	+	+	+	****	NADH dehydrogenase (ubiquinone) Fe-S protein 1, 75kDa (NADH-coenzyme Q reductase) precursor [Bos taurus]
5	gi 28461205	+	+	+	+	****	isocitrate dehydrogenase 2 (NADP+), mitochondrial [Bos taurus]
6	gi 28461221	+	+	+	+	****	ATP synthase, H+ transporting, mitochondrial F1 complex, beta subunit [Bos taurus]
7	gi 108809	+	+	+	-	***	NADH2 dehydrogenase (ubiquinone) (EC 1.6.5.3) 30K chain precursor - bovine
8	gi 117102	+	+	+	-	***	Cytochrome c oxidase polypeptide Vb (VI)
9	gi 27806307	+	+	+	-	***	mitochondrial ATP synthase, O subunit [Bos taurus]
10	gi 27806939	+	+	+	-	***	myoglobin [Bos taurus]
11	gi 27807305	+	+	+	-	***	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit d [Bos taurus]
12	gi 27819608	+	+	+	-	***	hemoglobin, beta [beta globin] [Bos taurus]
13	gi 284648	+	+	-	+	***	succinate dehydrogenase (ubiquinone) (EC 1.3.5.1) flavoprotein chain precursor, mitochondrial - bovine
14	gi 30794280	-	+	+	+	***	albumin [Bos taurus]
15	gi 40889866	+	+	+	-	***	Chain R, Bovine Heart Cytochrome C Oxidase At The Fully Reduced State
16	gi 437027	+	+	-	+	***	plasmalemmal porin (VDAC-1) [Bos taurus]
17	gi 51247981	+	+	-	+	***	Chain F, Beryllium Fluoride Inhibited Bovine F1-ATPase
18	gi 61813922	+	+	+	-	***	PREDICTED: similar to citrate (si)-synthase (EC 4.1.3.7) precursor - pig [Bos taurus]
19	gi 61820041	+	+	+	-	***	PREDICTED: similar to Acyl-CoA dehydrogenase, medium-chain specific, mitochondrial precursor (MCAD) [Bos taurus]
20	gi 61824969	+	+	+	-	***	PREDICTED: similar to Stress-70 protein, mitochondrial precursor (75 kDa glucose regulated protein) (GRP 75) (Peptide-binding protein 74) (PBP74) (Mortalin) (MOT) [Bos taurus]
21	gi 61867324	+	+	+	-	***	PREDICTED: similar to Malate dehydrogenase, mitochondrial precursor [Bos taurus]
22	gi 61882049	+	-	+	+	***	PREDICTED: similar to Creatine kinase, mitochondrial 2 [Bos taurus]
23	gi 833783	+	+	-	+	***	NADH:ubiquinone oxidoreductase (428 AA) [Bos taurus]
24	gi 114617	-	+	+	-	**	ATP synthase B chain, mitochondrial
25	gi 120604	+	+	-	-	**	Fumarate hydratase, mitochondrial (Fumarase)
26	gi 1352618	-	+	+	-	**	2-oxoglutarate dehydrogenase E1 component, mitochondrial precursor (Alpha-ketoglutarate dehydrogenase)
27	gi 1364245	+	-	+	-	**	NADH dehydrogenase 24 kDa subunit (AA 6-217) [Bos taurus]
28	gi 233598	+	-	+	-	**	13 kDa-B polypeptide of iron-sulfur protein fraction of NADH:ubiquinone oxidoreductase [cattle, heart, Peptide Mitochondrial Partial, 114 aa]
29	gi 27805909	-	+	-	+	**	dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit hydroxyacyl-Coenzyme A [Bos taurus]
30	gi 27806205	-	+	-	+	**	acyl-coenzyme A dehydrogenase, very long chain [Bos taurus]
31	gi 27806831	-	+	-	+	**	nicotinamide nucleotide transhydrogenase [Bos taurus]
32	gi 27807137	-	+	-	+	**	ubiquinol-cytochrome c reductase core protein I [Bos taurus]
33	gi 27807161	+	+	-	-	**	isocitrate dehydrogenase 3 (NAD+) alpha [Bos taurus]
34	gi 27807365	+	+	-	-	**	ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1 [Bos taurus]
35	gi 27807441	-	+	-	+	**	acetyl-CoA synthetase 2-like [Bos taurus]
36	gi 28461255	+	+	-	-	**	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 10, 22kDa [Bos taurus]
37	gi 28603782	+	-	+	-	**	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 10, 42kDa [Bos taurus]
38	gi 28603790	+	+	-	-	**	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7, 18kDa [Bos taurus]
39	gi 30583661	+	-	+	-	**	prohibitin [Homo sapiens]
40	gi 32189340	-	+	+	-	**	solute carrier family 25 member 4 [Bos taurus]

#	Accession	IEF-PAGE	Blue Native	Tricine-Urea	16-BAC	No. of Methods ^a	Protein
41	gi 34855983	-	+	-	+	**	"similar to inner membrane protein, mitochondrial (mitofilin)
42	gi 3851608	+	+	-	-	**	succinate dehydrogenase lp subunit [Bos taurus]
43	gi 40889865	-	+	+	-	**	Chain Q, Bovine Heart Cytochrome C Oxidase At The Fully Reduced State
44	gi 45430053	-	+	-	+	**	NADH dehydrogenase (ubiquinone) 42 kDa subunit [Bos taurus]
45	gi 4557409	-	-	+	+	**	cardiac calsequestrin 2 [Homo sapiens]
46	gi 4838363	+	+	-	-	**	creatine kinase M chain [Bos taurus]
47	gi 61553497	+	+	-	-	**	mitochondrial short-chain enoyl-coenzyme A hydratase 1 precursor [Bos taurus]
48	gi 61815075	+	-	+	-	**	PREDICTED: similar to 60 kDa heat shock protein, mitochondrial precursor (Hsp60) (60 kDa chaperonin) (CPN60) (Heat shock protein 60) (HSP-60) (Mitochondrial matrix protein P1) (P60 lymphocyte protein) (HuCHA60), partial [Bos taurus]
49	gi 61831678	-	+	-	+	**	PREDICTED: similar to Dihydrolipoamide dehydrogenase, precursor, partial [Bos taurus]
50	gi 61840143	-	+	+	-	**	PREDICTED: similar to prohibitin 2, partial [Bos taurus]
51	gi 61856478	+	+	-	-	**	PREDICTED: similar to cytosolic malate dehydrogenase [Bos taurus]
52	gi 61858852	+	+	-	-	**	PREDICTED: similar to mitochondrial acetoacetyl-CoA thiolase, partial [Bos taurus]
53	gi 61859294	+	+	-	-	**	PREDICTED: similar to 3-oxoacid CoA transferase 1, partial [Bos taurus]
54	gi 71621	+	-	-	+	**	actin beta - bovine (tentative sequence)
55	gi 71625	+	-	-	+	**	actin gamma - bovine (tentative sequence)
56	gi 1042206	+	-	-	-	*	purine nucleoside phosphorylase, PNP, purine nucleoside:orthophosphate ribosyltransferase {EC 2.4.2.1} [cattle, spleen, Peptide, 284 aa]
57	gi 1083063	+	-	-	-	*	protein disulfide-isomerase (EC 5.3.4.1) ER60 precursor - bovine
58	gi 109376	-	-	+	-	*	sarcoplasmic reticulum 53K glycoprotein precursor - rabbit
59	gi 117757	-	+	-	-	*	Cytochrome c1, heme protein, mitochondrial (Cytochrome c-1)
60	gi 125192	+	-	-	-	*	cAMP-dependent protein kinase type I-alpha regulatory subunit
61	gi 1351356	-	-	+	-	*	Ubiquinol-cytochrome c reductase complex 14 kDa protein (Complex III subunit VI)
62	gi 136691	+	-	-	-	*	Ubiquinol-cytochrome c reductase complex 11 kDa protein (Mitochondrial hinge protein) (Cytochrome C1, nonheme 11 kDa protein) (Complex III subunit VIII)
63	gi 14043738	-	-	-	+	*	CGI-51 protein [Homo sapiens]
64	gi 162648	+	-	-	-	*	albumin [Bos taurus]
65	gi 1703281	+	-	-	-	*	Cytosol aminopeptidase (Leucine aminopeptidase) (LAP) (Leucyl aminopeptidase) (Proline aminopeptidase) (Prolvl aminopeptidase)
66	gi 1943516	+	-	-	-	*	70kd Heat Shock Cognate Protein Atpase Domain, K71a Mutant
67	gi 21361103	-	-	-	+	*	solute carrier family 25 (mitochondrial carrier, Aralar), member 12 [Homo sapiens]
68	gi 2285903	+	-	-	-	*	glyceraldehyde 3-phosphate dehydrogenase [Bos taurus]
69	gi 233597	+	-	-	-	*	13 kDa-A polypeptide of iron-sulfur protein fraction of NADH:ubiquinone oxidoreductase [cattle, heart, Peptide Mitochondrial Partial, 96 aa]
70	gi 245563	+	-	-	-	*	apolipoprotein A-I, apoA-1 [Bos=cattle, Friesian-Holstein male calves aged 2-4 weeks, Peptide, 247 aa]
71	gi 2506441	-	+	-	-	*	Glyceraldehyde-3-phosphate dehydrogenase (GAPDH)
72	gi 266684	+	-	-	-	*	Dihydrolipoyllysine-residue succinyltransferase component of 2-oxoglutarate dehydrogenase complex mitochondrial precursor (Dihydrolipoamide succinyltransferase component of 2-oxoglutarate dehydrogenase complex) (E2) (E2K)
73	gi 27805849	+	-	-	-	*	crystallin, alpha polypeptide 2 [Bos taurus]
74	gi 27805907	+	-	-	-	*	hydroxyacyl-Coenzyme A dehydrogenase, type II hydroxyacyl-Coenzyme A [Bos taurus]
75	gi 27805917	+	-	-	-	*	histidine triad nucleotide binding protein 2 [Bos taurus]
76	gi 27805925	+	-	-	-	*	heat shock 70 kDa protein 8 [Bos taurus]
77	gi 27806319	+	-	-	-	*	adenylate kinase 3 [Bos taurus]
78	gi 27806367	+	-	-	-	*	Tu translation elongation factor, mitochondrial [Bos taurus]

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79	gi 27806561	+	-	-	-	*	lactate dehydrogenase B [Bos taurus]
80	gi 27806667	+	-	-	-	*	cytochrome b-5 [Bos taurus]
81	gi 27806785	+	-	-	-	*	vimentin [Bos taurus]
82	gi 27806929	+	-	-	-	*	adenylate kinase 2 [Bos taurus]
83	gi 27807167	+	-	-	-	*	anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2) [Bos taurus]
84	gi 27807263	-	-	+	-	*	tumor rejection antigen (gp96) 1 [Bos taurus]
85	gi 27807291	-	-	+	-	*	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F6 [Bos taurus]
86	gi 27807353	+	-	-	-	*	NADH dehydrogenase (ubiquinone) Fe-S protein 3, 30kDa (NADH-coenzyme Q reductase) [Bos taurus]
87	gi 27807359	+	-	-	-	*	NADH dehydrogenase (ubiquinone) Fe-S protein 8, 23kDa (NADH-coenzyme Q reductase) [Bos taurus]
88	gi 27807377	+	-	-	-	*	glutamic-oxaloacetic transaminase 2, mitochondrial (aspartate aminotransferase 2) [Bos taurus]
89	gi 27885005	-	+	-	-	*	dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta sub
90	gi 28189643	+	-	-	-	*	similar to B15 subunit of the NADH: ubiquinone oxidoreductase complex [Bos taurus]
91	gi 28461191	+	-	-	-	*	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 9, 22kDa [Bos taurus]
92	gi 28461207	+	-	-	-	*	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6, 14kDa [Bos taurus]
93	gi 28461257	-	-	+	-	*	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2, 8kDa [Bos taurus]
94	gi 28461259	-	-	+	-	*	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4, 9kDa [Bos taurus]
95	gi 28461263	-	+	-	-	*	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4, 15kDa [Bos taurus]
96	gi 28461275	+	-	-	-	*	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 8, 19kDa [Bos taurus]
97	gi 28461281	+	-	-	-	*	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5, 13kDa [Bos taurus]
98	gi 28603752	-	+	-	-	*	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit e [Bos taurus]
99	gi 28603758	-	+	-	-	*	NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2, 14.5kDa [Bos taurus]
100	gi 28603804	-	+	-	-	*	cell death-regulatory protein GRIM19 [Bos taurus]
101	gi 28603814	-	+	-	-	*	cytochrome c oxidase subunit VIb [Bos taurus]
102	gi 29126205	+	-	-	-	*	acetyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase) [Mus musculus]
103	gi 29135275	+	-	-	-	*	thiosulfate sulfurtransferase (rhodanese) [Bos taurus]
104	gi 29135329	+	-	-	-	*	glutathione S-transferase pi [Bos taurus]
105	gi 29351581	+	-	-	-	*	Electron transferring flavoprotein, beta polypeptide [Mus musculus]
106	gi 2959452	+	-	-	-	*	desmin [Bos taurus]
107	gi 29747871	-	-	-	+	*	Citrate synthase-like protein [Mus musculus]
108	gi 30794348	+	-	-	-	*	casein alpha-S1 [Bos taurus]
109	gi 31543274	+	-	-	-	*	metaxin 2 [Mus musculus]
110	gi 33357748	+	-	-	-	*	Chain H, Bovine Mitochondrial F1-ATPase Complexed With The Inhibitor Protein If1
111	gi 3660253	-	-	+	-	*	Chain F, The Structure Of Bovine F1-ATPase Covalently Inhibited With 4-Chloro-7-Nitrobenzofurazan
112	gi 387090	+	-	-	-	*	alpha-cardiac actin
113	gi 3891848	+	-	-	-	*	Chain A, Crystal Structure Of Bovine Mitochondrial Cytochrome Bc1 Complex, Alpha Carbon Atoms Only
114	gi 3891854	-	+	-	-	*	Chain G, Crystal Structure Of Bovine Mitochondrial Cytochrome Bc1 Complex, Alpha Carbon Atoms Only
115	gi 3915777	+	-	-	-	*	Dihydrolipoyllysine-residue acetyltransferase component of pyruvate dehydrogenase complex, mitochondrial precursor (Pyruvate dehydrogenase complex E2 subunit) (PDCE2) (E2) (Dihydrolipoamide S-acetyltransferase component of pyruvate dehydrogenase complex)
116	gi 39930503	+	-	-	-	*	ATP synthase, H+ transporting, mitochondrial F1 complex, gamma subunit [Rattus norvegicus]
117	gi 40889053	-	-	-	+	*	Chain R, Crystal Structure Of The Rabbit Muscle Glyceraldehyde-3- Phosphate Dehydrogenase (Gapdh)
118	gi 448581	+	-	-	-	*	pyruvate dehydrogenase:SUBUNIT=beta

#	Accession	IEF-PAGE	Blue Native	Tricine-Urea	16-BAC	No. of Methods ^a	Protein
119	gi 45430009	+	-	-	-	*	CGI-105 protein [Bos taurus]
120	gi 4558113	+	-	-	-	*	Chain A, Purine Nucleoside Phosphorylase
121	gi 48374083	+	-	-	-	*	annexin A4 [Bos taurus]
122	gi 498260	+	-	-	-	*	manganous superoxide dismutase
123	gi 51247188	-	+	-	-	*	Chain T, Bovine Cytochrome Bc1 Complex With Stigmatellin And Antimycin
124	gi 51338770	-	+	-	-	*	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial precursor (FP) (Flavoprotein subunit of complex II)
125	gi 52782253	-	-	-	+	*	oxoglutarate dehydrogenase [Macaca fascicularis]
126	gi 54310686	-	+	-	-	*	cytochrome c oxidase subunit II [Bos grunniens]
127	gi 543874	-	+	-	-	*	ATP synthase gamma chain, mitochondrial precursor
128	gi 55957259	+	-	-	-	*	succinate-CoA ligase, ADP-forming, beta subunit [Homo sapiens]
129	gi 57038366	+	-	-	-	*	PREDICTED: similar to heat shock protein, alpha-crystallin-related, B6 [Canis familiaris]
130	gi 57101266	+	-	-	-	*	PREDICTED: similar to Myosin light polypeptide 3 (Myosin light chain 1, slow-twitch muscle B/ventricular isoform) (MLC1SB) (Ventricular/slow twitch myosin alkali light chain) (Cardiac myosin light chain-1) (CMLC1) [Canis familiaris]
131	gi 57208844	-	-	-	+	*	programmed cell death 8 (apoptosis-inducing factor) [Homo sapiens]
132	gi 57209621	+	-	-	-	*	pyruvate dehydrogenase (lipoamide) alpha 1 [Homo sapiens]
133	gi 57657	+	-	-	-	*	pyruvate dehydrogenase E1 alpha form 1 subunit [Rattus rattus]
134	gi 59857635	-	+	-	-	*	acyl-Coenzyme A dehydrogenase, short/branched chain precursor [Bos taurus]
135	gi 59858339	+	-	-	-	*	coiled-coil-helix-coiled-coil-helix domain containing 3 [Bos taurus]
136	gi 59858341	+	-	-	-	*	superoxide dismutase 2, mitochondrial [Bos taurus]
137	gi 59858383	-	+	-	-	*	lactate dehydrogenase B [Bos taurus]
138	gi 6006421	+	-	-	-	*	hemoglobin alpha chain [Bos taurus]
139	gi 61553356	-	+	-	-	*	purine nucleoside phosphorylase [Bos taurus]
140	gi 61553952	-	-	+	-	*	retinol binding protein 1, cellular [Bos taurus]
141	gi 61554381	-	+	-	-	*	isocitrate dehydrogenase 3 (NAD+) gamma isoform a precursor [Bos taurus]
142	gi 61554470	+	-	-	-	*	isocitrate dehydrogenase 3 (NAD+) gamma isoform a precursor - bovine
143	gi 61813387	+	-	-	-	*	PREDICTED: similar to ATPase, aminophospholipid transporter-like, class I, type 8A, member 2, partial [Bos taurus]
144	gi 61814899	-	+	-	-	*	PREDICTED: similar to Carnitine palmitoyltransferase II, partial [Bos taurus]
145	gi 61817069	+	-	-	-	*	PREDICTED: similar to mitochondrial ribosomal protein L50 [Bos taurus]
146	gi 61822295	+	-	-	-	*	PREDICTED: similar to ES1 protein homolog, mitochondrial precursor (Protein KNP-I) (GT335 protein), partial [Bos taurus]
147	gi 61828933	+	-	-	-	*	PREDICTED: similar to Pyruvate dehydrogenase protein X component, mitochondrial precursor (Dihydrolipoamide dehydrogenase-binding protein of pyruvate dehydrogenase complex) (Lipoal-containing pyruvate dehydrogenase complex component X) (E3-binding prote
148	gi 61830576	+	-	-	-	*	PREDICTED: similar to E-1 beta subunit of the pyruvate dehydrogenase complex, partial [Bos taurus]
149	gi 61831186	+	-	-	-	*	PREDICTED: similar to acyl-CoA dehydrogenase (EC 1.3.99.3) precursor, short-chain-specific - rat, partial [Bos taurus]
150	gi 61831254	-	+	-	-	*	PREDICTED: similar to Short chain 3-hydroxyacyl-CoA dehydrogenase, mitochondrial precursor (HCDH) (Medium and short chain L-3-hydroxyacyl-coenzyme A dehydrogenase), partial [Bos taurus]
151	gi 61835358	+	-	-	-	*	PREDICTED: similar to Beta enolase (2-phospho-D-glycerate hydro-lyase) (Muscle-specific enolase) (MSE) (Skeletal muscle enolase) (Enolase 3), partial [Bos taurus]
152	gi 61835610	-	-	+	-	*	PREDICTED: similar to mitofilin, partial [Bos taurus]
153	gi 61838105	+	-	-	-	*	PREDICTED: similar to Short chain 3-hydroxyacyl-CoA dehydrogenase, mitochondrial precursor (HCDH) (Medium and short chain L-3-hydroxyacyl-coenzyme A dehydrogenase), partial [Bos taurus]
154	gi 61841483	-	-	+	-	*	PREDICTED: similar to actinin alpha 2, partial [Bos taurus]
155	gi 61842751	+	-	-	-	*	PREDICTED: similar to serine/threonine/tyrosine kinase 1, partial [Bos taurus]

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156	gi 61844553	-	+	-	-	*	PREDICTED: similar to 60 kDa heat shock protein, mitochondrial precursor (Hsp60) (60 kDa chaperonin) (CPN60) (Heat shock protein 60) (HSP-60) (Mitochondrial matrix protein P1) (P60 lymphocyte protein) (HuCHA60), partial [Bos taurus]
157	gi 61850233	+	-	-	-	*	PREDICTED: similar to hypothetical protein, partial [Bos taurus]
158	gi 61856242	+	-	-	-	*	PREDICTED: similar to succinyl-CoA synthetase beta-subunit, partial [Bos taurus]
159	gi 61858132	-	+	-	-	*	PREDICTED: similar to Alpha-2-macroglobulin precursor (Alpha-2-M), partial [Bos taurus]
160	gi 61861654	-	+	-	-	*	PREDICTED: similar to valosin precursor [Bos taurus]
161	gi 61863144	+	-	-	-	*	PREDICTED: similar to FabG-like protein, partial [Bos taurus]
162	gi 61866326	+	-	-	-	*	PREDICTED: similar to myosin light chain 2, partial [Bos taurus]
163	gi 61866502	-	+	-	-	*	PREDICTED: similar to ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g [Bos taurus]
164	gi 61870138	+	-	-	-	*	PREDICTED: similar to 3-hydroxybutyrate dehydrogenase precursor [Bos taurus]
165	gi 61874440	+	-	-	-	*	PREDICTED: similar to phosphoglycerate mutase 1 (brain) [Bos taurus]
166	gi 61876588	+	-	-	-	*	PREDICTED: similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) [Bos taurus]
167	gi 61885153	+	-	-	-	*	PREDICTED: similar to Hsp27, partial [Bos taurus]
168	gi 61886457	+	-	-	-	*	PREDICTED: similar to Ribosomal protein S6 kinase alpha 6 (S6K-alpha 6) (90 kDa ribosomal protein S6 kinase 6) (p90-RSK 6) (Ribosomal S6 kinase 4) (RSK-4) (pp90RSK4), partial [Bos taurus]
169	gi 61887001	+	-	-	-	*	PREDICTED: similar to QIL1 protein [Bos taurus]
170	gi 61888856	+	-	-	-	*	triosephosphate isomerase [Bos taurus]
171	gi 627764	+	-	-	-	*	substrate protein of mitochondrial ATP-dependent proteinase SP-22 - bovine
172	gi 6729922	+	-	-	-	*	Chain A, Nmr Study Of Bovine Heart Fatty Acid Binding Protein
173	gi 8574295	+	-	-	-	*	voltage-dependent anion channel 2 [Bos taurus]
174	gi 9651188	+	-	-	-	*	cardiac myosin light chain-1 [Homo sapiens]

^a The number of asteriks denotes the number of methods in which the respective protein has been identified