

Protein Ratio Feature Table		
Master N	Accession	Name
1	sp Q28343 PGCA_CANFA	Aggrecan core protein OS=Canis familiaris GN=ACAN PE=2 SV=2
2	sp P07589 FINC_BOVIN	Fibronectin OS=Bos taurus GN=FN1 PE=1 SV=4
3	sp P08670 VIME_HUMAN	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4
4	sp Q13813-2 SPTA2_HUMAN	Isoform 2 of Spectrin alpha chain, brain OS=Homo sapiens GN=SPTAN1
5	sp P17182 ENOA_MOUSE	Alpha-enolase OS=Mus musculus GN=Eno1 PE=1 SV=3
6	sp P08728 K1C19_BOVIN	Keratin, type I cytoskeletal 19 OS=Bos taurus GN=KRT19 PE=2 SV=1
7	sp P25473 CLUS_CANFA	Clusterin OS=Canis familiaris GN=CLU PE=2 SV=1
8	sp P05787 K2C8_HUMAN	Keratin, type II cytoskeletal 8 OS=Homo sapiens GN=KRT8 PE=1 SV=7
9	sp Q4KY93 G3P_SPECI	Glyceraldehyde-3-phosphate dehydrogenase OS=Spermophilus citellus GN=GAPDH PE=2 SV=3
10	sp P14618-2 KPYM_HUMAN	Isoform M1 of Pyruvate kinase isozymes M1/M2 OS=Homo sapiens GN=PKM2
11	sp Q15335 CHAD_HUMAN	Chondroadherin OS=Homo sapiens GN=CHAD PE=2 SV=2
12	sp Q62261 SPTB2_MOUSE	Spectrin beta chain, brain 1 OS=Mus musculus GN=Sptbn1 PE=1 SV=2
13	sp Q7YQC6 HSP71_CANFA	Heat shock 70 kDa protein 1 OS=Canis familiaris GN=HSPA1 PE=2 SV=1
14	sp Q9NZN4 EHD2_HUMAN	EH domain-containing protein 2 OS=Homo sapiens GN=EHD2 PE=1 SV=2
15	sp Q3T0P6 PGK1_BOVIN	Phosphoglycerate kinase 1 OS=Bos taurus GN=PGK1 PE=2 SV=3
16	sp P12111 CO6A3_HUMAN	Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 PE=1 SV=4
17	sp Q29393 PGS2_CANFA	Decorin OS=Canis familiaris GN=DCN PE=2 SV=2
18	sp A5D7D1 ACTN4_BOVIN	Alpha-actinin-4 OS=Bos taurus GN=ACTN4 PE=2 SV=1
19	sp P63261 ACTG_HUMAN	Actin, cytoplasmic 2 OS=Homo sapiens GN=ACTG1 PE=1 SV=1
20	sp P02458 CO2A1_HUMAN	Collagen alpha-1(II) chain OS=Homo sapiens GN=COL2A1 PE=1 SV=3
21	sp Q5OQ56 XYLT1_CANFA	Xylyltransferase 1 OS=Canis familiaris GN=XYLT1 PE=2 SV=1
22	sp Q09666 AHNK_HUMAN	Neuroblast differentiation-associated protein AHNK OS=Homo sapiens GN=AHNAK PE=1 SV=2
23	sp P54714 TPIS_CANFA	Triosephosphate isomerase OS=Canis familiaris GN=TP1I PE=1 SV=3
24	sp Q6TEQ7 ANXA2_CANFA	Annexin A2 OS=Canis familiaris GN=ANXA2 PE=1 SV=1
25	sp A1A4K5 ENPP2_BOVIN	Ectonucleotide pyrophosphatase/phosphodiesterase family member 2 OS=Bos taurus GN=ENPP2 PE=1 SV=1
26	sp P16015 CAH3_MOUSE	Carbonic anhydrase 3 OS=Mus musculus GN=Ca3 PE=1 SV=3
27	sp Q9NZN3 EHD3_HUMAN	EH domain-containing protein 3 OS=Homo sapiens GN=EHD3 PE=1 SV=2
28	sp Q684U9 PRDX1_MYOLU	Peroxiredoxin-1 OS=Myotis lucifugus GN=PRDX1 PE=1 SV=1
29	sp Q9DBU1 PGAM1_MOUSE	Phosphoglycerate mutase 1 OS=Mus musculus GN=Pgam1 PE=1 SV=3
30	sp Q99985 SEM3C_HUMAN	Semaphorin-3C OS=Homo sapiens GN=SEMA3C PE=1 SV=2
31	sp P35444 COMP_RAT	Cartilage oligomeric matrix protein OS=Rattus norvegicus GN=Comp PE=1 SV=1
32	sp P30427 PLEC_RAT	Plectin OS=Rattus norvegicus GN=Plec PE=1 SV=2
33	sp Q4R586 LDHB_MACFA	L-lactate dehydrogenase B chain OS=Macaca fascicularis GN=LDHB PE=2 SV=3
34	sp P04075 ALDOA_HUMAN	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2
35	sp Q258K2 MYH9_CANFA	Myosin-9 OS=Canis familiaris GN=MYH9 PE=2 SV=1
36	sp P05784 K1C18_MOUSE	Keratin, type I cytoskeletal 18 OS=Mus musculus GN=Krt18 PE=1 SV=5
37	sp P26038 MOES_HUMAN	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3
38	sp P62936 PIIA_PIG	Peptidyl-prolyl cis-trans isomerase A OS=Sus scrofa GN=PIIA PE=1 SV=2
39	sp P47942 DPYL2_RAT	Dihydropyrimidinase-related protein 2 OS=Rattus norvegicus GN=Dpysl2 PE=1 SV=1
40	sp Q28275 FINC_CANFA	Fibronectin (Fragment) OS=Canis familiaris GN=FN1 PE=2 SV=2
41	sp Q66K08 CILP1_MOUSE	Cartilage intermediate layer protein 1 OS=Mus musculus GN=Cilp PE=2 SV=1
42	sp Q6NZ12 PTRF_HUMAN	Polymerase I and transcript release factor OS=Homo sapiens GN=PTRF PE=1 SV=1
43	sp Q75094 SLIT3_HUMAN	Slit homolog 3 protein OS=Homo sapiens GN=SLIT3 PE=2 SV=2
44	sp Q6P9T8 TBB2C_RAT	Tubulin beta-2C chain OS=Rattus norvegicus GN=Tubb2c PE=1 SV=1
45	sp P19858 LDHA_BOVIN	L-lactate dehydrogenase A chain OS=Bos taurus GN=LDHA PE=2 SV=2
46	sp Q75369 FLNB_HUMAN	Filamin-B OS=Homo sapiens GN=FLNB PE=1 SV=2
47	sp P08059 G6PI_PIG	Glucose-6-phosphate isomerase OS=Sus scrofa GN=GPI PE=1 SV=3
48	sp Q727G0 TARSH_HUMAN	Target of Nesh-SH3 OS=Homo sapiens GN=ABI3BP PE=1 SV=1
49	sp P02751 FINC_HUMAN	Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4
50	sp Q16851 UGPA_HUMAN	UTP--glucose-1-phosphate uridylyltransferase OS=Homo sapiens GN=UGP2 PE=1 SV=5
51	sp Q28381 HPLN1_HORSE	Hyaluronan and proteoglycan link protein 1 OS=Equus caballus GN=HAPLN1 PE=2 SV=1
52	sp Q71U36 TBA1A_HUMAN	Tubulin alpha-1A chain OS=Homo sapiens GN=TUBA1A PE=1 SV=1
53	sp P10412 H14_HUMAN	Histone H1.4 OS=Homo sapiens GN=HIST1H1E PE=1 SV=2
54	sp P18649 APOE_CANFA	Apolipoprotein E OS=Canis familiaris GN=APOE PE=1 SV=2
55	sp Q66RN5 EF1A1_FELCA	Elongation factor 1-alpha 1 OS=Felis catus GN=EEF1A1 PE=2 SV=1
56	sp Q02678 PGS1_CANFA	Biglycan OS=Canis familiaris GN=BGN PE=2 SV=1
57	sp Q10758 K2C8_RAT	Keratin, type II cytoskeletal 8 OS=Rattus norvegicus GN=Krt8 PE=1 SV=3
58	sp Q71U34 HSP7C_SAGOE	Heat shock cognate 71 kDa protein OS=Saguinus oedipus GN=HSPA8 PE=2 SV=1
59	sp P05124 KCRB_CANFA	Creatine kinase B-type OS=Canis familiaris GN=CKB PE=1 SV=1
60	sp Q2PFV7 ACTN1_MACFA	Alpha-actinin-1 OS=Macaca fascicularis GN=ACTN1 PE=2 SV=1
61	sp P49822 ALBU_CANFA	Serum albumin OS=Canis familiaris GN=ALB PE=1 SV=3
62	sp P31404 VATA_BOVIN	V-type proton ATPase catalytic subunit A OS=Bos taurus GN=ATP6V1A PE=2 SV=2
63	sp P23528 COF1_HUMAN	Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3
64	sp P29361 14332_SHEEP	14-3-3 protein zeta/delta OS=Ovis aries GN=YWHAZ PE=1 SV=1
65	sp P12109 CO6A1_HUMAN	Collagen alpha-1(VI) chain OS=Homo sapiens GN=COL6A1 PE=1 SV=3
66	sp Q68B18 OLML2B_HUMAN	Olfactomedin-like protein 2B OS=Homo sapiens GN=OLFML2B PE=2 SV=2
67	sp P07900-2 HSP90A_HUMAN	Isoform HSP90AA1-2 of Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1
68	sp P38486 LEG3_CANFA	Galectin-3 OS=Canis familiaris GN=LGALS3 PE=1 SV=3
69	sp Q63279 K1C19_RAT	Keratin, type I cytoskeletal 19 OS=Rattus norvegicus GN=Krt19 PE=1 SV=2
70	sp P48678 LMNA_MOUSE	Lamin-A/C OS=Mus musculus GN=Lmna PE=1 SV=2

71	sp P98160 PGBM_HUMAN	Basement membrane-specific heparan sulfate proteoglycan core protein OS=Homo sapiens GN=HSPG2 PE=1 SV=3
72	sp P42929 HSPB1_CANFA	Heat shock protein beta-1 OS=Canis familiaris GN=HSPB1 PE=2 SV=1
73	sp P60526 HBB_CHRRB	Hemoglobin subunit beta OS=Chrysocony brachyurus GN=HBB PE=1 SV=1
74	sp P18206 VINC_HUMAN	Vinculin OS=Homo sapiens GN=VCL PE=1 SV=4
75	sp Q3Y1X4 PEBP1_CANFA	Phosphatidylethanolamine-binding protein 1 OS=Canis familiaris GN=PEBP1 PE=1 SV=1
76	sp P47856 GFPT1_MOUSE	Glucosamine-fructose-6-phosphate aminotransferase [isomerizing] 1 OS=Mus musculus GN=Gfpt1 PE=1 SV=3
77	sp P02510 CRYAB_BOVIN	Alpha-crystallin B chain OS=Bos taurus GN=CRYAB PE=1 SV=2
78	sp Q9R118 HTRA1_MOUSE	Serine protease HTRA1 OS=Mus musculus GN=Htra1 PE=2 SV=1
79	sp Q77834 PRDX6_BOVIN	Peroxiredoxin-6 OS=Bos taurus GN=PRDX6 PE=1 SV=3
80	sp P21333 FLNA_HUMAN	Filamin-A OS=Homo sapiens GN=FLNA PE=1 SV=4
81	sp Q9EPW4 CLC3A_MOUSE	C-type lectin domain family 3 member A OS=Mus musculus GN=Clec3a PE=2 SV=1
82	sp P51884 LUM_HUMAN	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2
83	sp Q9J1M9-2 PRG4_MOUSE	Isoform B of Proteoglycan 4 OS=Mus musculus GN=PrG4
84	sp Q32524 COBA2_BOVIN	Collagen alpha-2(XI) chain OS=Bos taurus GN=COL11A2 PE=3 SV=1
85	sp P60530 HBA_CANLA	Hemoglobin subunit alpha OS=Canis latrans GN=HBA PE=1 SV=2
86	sp P81709 LYSC2_CANFA	Lysozyme C, spleen isozyme OS=Canis familiaris PE=1 SV=1
87	sp P13608 PGCA_BOVIN	Aggrecan core protein OS=Bos taurus GN=ACAN PE=1 SV=3
88	sp P53396 ACLY_HUMAN	ATP-citrate synthase OS=Homo sapiens GN=ACLY PE=1 SV=3
89	sp Q3ZB28 STIP1_BOVIN	Stress-induced-phosphoprotein 1 OS=Bos taurus GN=STIP1 PE=2 SV=1
90	sp P12110 CO6A2_HUMAN	Collagen alpha-2(VI) chain OS=Homo sapiens GN=COL6A2 PE=1 SV=4
91	sp Q5R823 EF2_PONAB	Elongation factor 2 OS=Pongo abelii GN=EEF2 PE=2 SV=3
92	sp P00918 CAH2_HUMAN	Carbonic anhydrase 2 OS=Homo sapiens GN=CA2 PE=1 SV=2
93	sp P07093 GDN_HUMAN	Glia-derived nexin OS=Homo sapiens GN=SERPINE2 PE=1 SV=1
94	sp Q64610 ENPP2_RAT	Ectonucleotide pyrophosphatase/phosphodiesterase family member 2 OS=Rattus norvegicus GN=Enpp2 PE=1 SV=2
95	sp Q99497 PARK7_HUMAN	Protein DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2
96	sp Q6T308 GLGB_FELCA	1,4-alpha-glucan-branching enzyme OS=Felis catus GN=GBE1 PE=2 SV=1
97	sp Q9H4F8-2 SMOC1_HUMAN	Isoform 2 of SPARC-related modular calcium-binding protein 1 OS=Homo sapiens GN=SMOC1
98	sp P62992 RS27A_BOVIN	Ubiquitin-40S ribosomal protein S27a OS=Bos taurus GN=RS27A PE=1 SV=2
99	sp P00761 TRYP_PIG	Trypsin OS=Sus scrofa PE=1 SV=1
100	sp Q9GKN8 PRELP_BOVIN	Prolargin OS=Bos taurus GN=PRELP PE=2 SV=1
101	sp Q00394 A1AT_CERAE	Alpha-1-antitrypsin (Fragment) OS=Cercopithecus aethiops GN=SERPINA1 PE=2 SV=1
102	sp Q7YQD7 FBLN3_MACFA	EGF-containing fibulin-like extracellular matrix protein 1 OS=Macaca fascicularis GN=EFEMP1 PE=2 SV=1
103	sp P19803 GDIR1_BOVIN	Rho GDP-dissociation inhibitor 1 OS=Bos taurus GN=ARHGDI1 PE=1 SV=3
104	sp P06396 GELS_HUMAN	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1
105	sp P58771-2 TPM1_MOUSE	Isoform Fibroblast of Tropomyosin alpha-1 chain OS=Mus musculus GN=Tpm1
106	sp Q5R7K9 EDIL3_PONAB	EGF-like repeat and discoidin 1-like domain-containing protein 3 OS=Pongo abelii GN=EDIL3 PE=2 SV=1
107	sp P50609 FMOD_RAT	Fibromodulin OS=Rattus norvegicus GN=Fmod PE=2 SV=1
108	sp Q7YRU4 MDHC_FELCA	Malate dehydrogenase, cytoplasmic OS=Felis catus GN=MDH1 PE=2 SV=3
109	sp P11980-2 KPYM_RAT	Isoform M2 of Pyruvate kinase isozymes M1/M2 OS=Rattus norvegicus GN=Pkm2
110	sp Q0VCX2 GRP78_BOVIN	78 kDa glucose-regulated protein OS=Bos taurus GN=HSPA5 PE=2 SV=1
111	sp Q97555 GDIA_CANFA	Rab GDP dissociation inhibitor alpha OS=Canis familiaris GN=GDII1 PE=2 SV=1
112	sp Q92954-2 PRG4_HUMAN	Isoform B of Proteoglycan 4 OS=Homo sapiens GN=PRG4
113	sp Q99536 VAT1_HUMAN	Synaptic vesicle membrane protein VAT-1 homolog OS=Homo sapiens GN=VAT1 PE=1 SV=2
114	sp P62262 1433E_SHEEP	14-3-3 protein epsilon OS=Ovis aries GN=YWHA6 PE=1 SV=1
115	sp Q68FD5 CLH_MOUSE	Clathrin heavy chain 1 OS=Mus musculus GN=Cltc PE=1 SV=3
116	sp Q5R4B4 NP11_PONAB	Nucleosome assembly protein 1-like 1 OS=Pongo abelii GN=NAP11L1 PE=2 SV=1
117	sp P24821 TENA_HUMAN	Tenascin OS=Homo sapiens GN=TNC PE=1 SV=3
118	sp P23352 KALM_HUMAN	Anosmin-1 OS=Homo sapiens GN=KAL1 PE=1 SV=3
119	sp Q63355 MYO1C_RAT	Myosin-1c OS=Rattus norvegicus GN=Myo1c PE=2 SV=2
120	sp Q9XSA7 CLIC4_BOVIN	Chloride intracellular channel protein 4 OS=Bos taurus GN=CLIC4 PE=2 SV=3
121	sp P12107-2 COBA1_HUMAN	Isoform B of Collagen alpha-1(XI) chain OS=Homo sapiens GN=COL11A1
122	sp Q50KA8 NDKB_CANFA	Nucleoside diphosphate kinase B OS=Canis familiaris GN=NME2 PE=2 SV=1
123	sp P79385 MFGM_PIG	Lactadherin OS=Sus scrofa GN=MFG6E PE=1 SV=2
124	sp P52907 CAZA1_HUMAN	F-actin-capping protein subunit alpha-1 OS=Homo sapiens GN=CAPZA1 PE=1 SV=3
125	sp P53534 PYGB_RAT	Glycogen phosphorylase, brain form (Fragment) OS=Rattus norvegicus GN=Pygb PE=1 SV=3
126	sp Q5R9X1 CADH2_PONAB	Cadherin-2 OS=Pongo abelii GN=CDH2 PE=2 SV=1
127	sp Q8R1M2 H2AJ_MOUSE	Histone H2A.J OS=Mus musculus GN=H2afj PE=2 SV=1
128	sp A3KMV5 UBA1_BOVIN	Ubiquitin-like modifier-activating enzyme 1 OS=Bos taurus GN=UBA1 PE=2 SV=1
129	sp Q5RCP8 H2B2E_PONAB	Histone H2B type 2-E OS=Pongo abelii GN=HIST2H2BE PE=2 SV=3
130	sp P31836 NCAM1_BOVIN	Neural cell adhesion molecule 1 OS=Bos taurus GN=NCAM1 PE=1 SV=1
131	sp P01023 AZMG_HUMAN	Alpha-2-macroglobulin OS=Homo sapiens GN=AZM PE=1 SV=2
132	sp P52209 PGD_HUMAN	6-phosphogluconate dehydrogenase, decarboxylating OS=Homo sapiens GN=PGD PE=1 SV=3
133	sp P48616 VIME_BOVIN	Vimentin OS=Bos taurus GN=VIM PE=1 SV=3
134	sp Q8IUJ8 CILP2_HUMAN	Cartilage intermediate layer protein 2 OS=Homo sapiens GN=CILP2 PE=2 SV=2
135	sp Q9UKZ9 PCOC2_HUMAN	Procollagen C-endopeptidase enhancer 2 OS=Homo sapiens GN=PCOLCE2 PE=1 SV=1
136	sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1
137	sp P33724 CAV1_CANFA	Caveolin-1 OS=Canis familiaris GN=CAV1 PE=1 SV=1
138	sp P51662 ANXA1_RABIT	Annexin A1 OS=Oryctolagus cuniculus GN=ANXA1 PE=1 SV=1
139	sp Q5RCS7 H4_PONAB	Histone H4 OS=Pongo abelii PE=3 SV=1
140	sp Q3ZBT1 TERA_BOVIN	Transitional endoplasmic reticulum ATPase OS=Bos taurus GN=VCP PE=1 SV=1
141	sp P05786 K2C8_BOVIN	Keratin, type II cytoskeletal 8 OS=Bos taurus GN=KRT8 PE=2 SV=3
142	sp P35445 COMP_BOVIN	Cartilage oligomeric matrix protein OS=Bos taurus GN=COMP PE=1 SV=2
143	sp P17183 ENOG_MOUSE	Gamma-enolase OS=Mus musculus GN=Eno2 PE=1 SV=2
144	sp Q46379 LUM_RABIT	Lumican (Fragment) OS=Oryctolagus cuniculus GN=LUM PE=2 SV=1
145	sp Q5R1X4 ALDOC_PANTR	Fructose-bisphosphate aldolase C OS=Pan troglodytes GN=ALDOC PE=2 SV=3
146	sp Q9TS87 TAGL_BOVIN	Transgelin OS=Bos taurus GN=TAGLN PE=1 SV=4
147	sp Q62644 LECT2_BOVIN	Leukocyte cell-derived chemotaxin-2 OS=Bos taurus GN=LECT2 PE=1 SV=1
148	sp Q7LGC8 CHST3_HUMAN	Carbohydrate sulfotransferase 3 OS=Homo sapiens GN=CHST3 PE=1 SV=3
149	sp Q6YNX6 CALM_SHEEP	Calmodulin OS=Ovis aries GN=CALM2 PE=2 SV=3
150	sp P53601-2 A4_MACFA	Isoform APP695 of Amyloid beta A4 protein OS=Macaca fascicularis GN=APP
151	sp Q8HZQ5 EZRI_RABIT	Ezrin OS=Oryctolagus cuniculus GN=EZR PE=1 SV=3
152	sp P80895 PIMT_PIG	Protein-L-isoaspartate(D-aspartate) O-methyltransferase OS=Sus scrofa GN=PCMT1 PE=1 SV=3
153	sp Q13642-1 FHL1_HUMAN	Isoform SLIM1 of Four and a half LIM domains protein 1 OS=Homo sapiens GN=FHL1
154	sp P54652 HSP72_HUMAN	Heat shock-related 70 kDa protein 2 OS=Homo sapiens GN=HSPA2 PE=1 SV=1
155	sp P62815 VATB2_RAT	V-type proton ATPase subunit B, brain isoform OS=Rattus norvegicus GN=Atp6v1b2 PE=1 SV=1
156	sp Q5IS74 TPP1_PANTR	Tripeptidyl-peptidase 1 OS=Pan troglodytes GN=TPP1 PE=2 SV=1
157	sp Q64119 MYL6_RAT	Myosin light polypeptide 6 OS=Rattus norvegicus GN=Myf6 PE=1 SV=3
158	sp Q710C4 SAHH_PIG	Adenosylhomocysteinease OS=Sus scrofa GN=AHCY PE=3 SV=3
159	sp P50994 ANXA4_CANFA	Annexin A4 OS=Canis familiaris GN=ANXA4 PE=1 SV=2
160	sp Q55226 CHAD_MOUSE	Chondroadherin OS=Mus musculus GN=Chad PE=2 SV=1

161	sp P14141 CAH3_RAT	Carbonic anhydrase 3 OS=Rattus norvegicus GN=Ca3 PE=1 SV=3
162	sp Q8WNN6 SODC_CANFA	Superoxide dismutase [Cu-Zn] OS=Canis familiaris GN=SOD1 PE=2 SV=1
163	sp Q13126 MTAP_HUMAN	S-methyl-5'-thioadenosine phosphorylase OS=Homo sapiens GN=MTAP PE=1 SV=2
164	sp Q5R859 K1C19_PONAB	Keratin, type I cytoskeletal 19 OS=Pongo abelii GN=KRT19 PE=2 SV=1
165	sp P11276 FINC_MOUSE	Fibronectin OS=Mus musculus GN=Fn1 PE=1 SV=3
166	sp P05783 K1C18_HUMAN	Keratin, type I cytoskeletal 18 OS=Homo sapiens GN=KRT18 PE=1 SV=2
167	sp Q2PF23 PRDX2_MACFA	Peroxiredoxin-2 OS=Macaca fascicularis GN=PRDX2 PE=2 SV=3
168	sp Q5R8R1 ARP3_PONAB	Actin-related protein 3 OS=Pongo abelii GN=ACTR3 PE=2 SV=3
169	sp P56480 ATPB_MOUSE	ATP synthase subunit beta, mitochondrial OS=Mus musculus GN=Atp5b PE=1 SV=2
170	sp Q8HZQ0 RNAS4_PANTR	Ribonuclease 4 OS=Pan troglodytes GN=RNASE4 PE=3 SV=1
171	sp Q5REN2 PSA1_PONAB	Proteasome subunit alpha type-1 OS=Pongo abelii GN=PSMA1 PE=2 SV=1
172	sp Q28008 CLC3A_BOVIN	C-type lectin domain family 3 member A OS=Bos taurus GN=CLEC3A PE=2 SV=1
173	sp Q9H299 SH3L3_HUMAN	SH3 domain-binding glutamic acid-rich-like protein 3 OS=Homo sapiens GN=SH3BGL3 PE=1 SV=1
174	sp P076041 NEBL_HUMAN	Nebulette OS=Homo sapiens GN=NEBL PE=2 SV=1
175	sp Q9Y490 TLN1_HUMAN	Talin-1 OS=Homo sapiens GN=TLN1 PE=1 SV=3
176	sp Q5R4C1 TKT_PONAB	Transketolase OS=Pongo abelii GN=TKT PE=2 SV=1
177	sp A1A41 K6PL_BOVIN	6-phosphofructokinase, liver type OS=Bos taurus GN=PFKL PE=2 SV=1
178	sp A507R1 TR11B_BOVIN	Tumor necrosis factor receptor superfamily member 11B OS=Bos taurus GN=TNFRSF11B PE=2 SV=1
179	sp P13697 MAOX_RAT	NADP-dependent malic enzyme OS=Rattus norvegicus GN=Me1 PE=1 SV=2
180	sp P02459 CO2A1_BOVIN	Collagen alpha-1(I) chain (Fragments) OS=Bos taurus GN=COL2A1 PE=1 SV=3
181	sp P13942 COBA2_HUMAN	Collagen alpha-2(XI) chain OS=Homo sapiens GN=COL11A2 PE=1 SV=4
182	sp P80928 MIF_PIG	Macrophage migration inhibitory factor OS=Sus scrofa GN=MIF PE=1 SV=3
183	sp Q4R5F2 LGLU_MACFA	Lactylglutathione lyase OS=Macaca fascicularis GN=GLO1 PE=2 SV=3
184	sp P08729 K2C7_HUMAN	Keratin, type II cytoskeletal 7 OS=Homo sapiens GN=KRT7 PE=1 SV=4
185	sp Q2TBQ8 6PGL_BOVIN	6-phosphogluconolactonase OS=Bos taurus GN=PGLS PE=2 SV=1
186	sp Q6VHK3 CD109_HUMAN	CD109 antigen OS=Homo sapiens GN=CD109 PE=1 SV=2
187	sp Q9QP5 PRELP_RAT	Prolargin OS=Rattus norvegicus GN=Prelp PE=2 SV=1
188	sp Q6RUV5 RAC1_RAT	Ras-related C3 botulinum toxin substrate 1 OS=Rattus norvegicus GN=Rac1 PE=1 SV=1
189	sp Q95340 PAPS2_HUMAN	Bifunctional 3'-phosphoadenosine 5'-phosphosulfate synthase 2 OS=Homo sapiens GN=PAPSS2 PE=1 SV=2
190	sp P20112 SPRC_PIG	SPARC OS=Sus scrofa GN=SPARC PE=1 SV=2
191	sp Q91V12 PRDBP_MOUSE	Protein kinase C delta-binding protein OS=Mus musculus GN=Prckdbp PE=1 SV=1
192	sp Q7JUJ6 TBB5_PANTR	Tubulin beta chain OS=Pan troglodytes GN=TUBB PE=3 SV=1
193	sp P20618 PSB1_HUMAN	Proteasome subunit beta type-1 OS=Homo sapiens GN=PSMB1 PE=1 SV=2
194	sp Q7SIB7 PGK1_PIG	Phosphoglycerate kinase 1 OS=Sus scrofa GN=PGK1 PE=1 SV=3
195	sp Q5R1W0 ANXA5_PANTR	Annexin A5 OS=Pan troglodytes GN=ANXA5 PE=2 SV=3
196	sp P06733 ENO4_HUMAN	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2
197	sp P07451 CAH3_HUMAN	Carbonic anhydrase 3 OS=Homo sapiens GN=CA3 PE=1 SV=3
198	sp Q61792 LASP1_MOUSE	LIM and SH3 domain protein 1 OS=Mus musculus GN=Lasp1 PE=1 SV=1
199	sp P05114 HMGN1_HUMAN	Non-histone chromosomal protein HMG-14 OS=Homo sapiens GN=HMGN1 PE=1 SV=3
200	sp Q5SRC20 1433G_PONAB	14-3-3 protein gamma OS=Pongo abelii GN=YWHAG PE=2 SV=3
201	sp Q66H80 COPD_RAT	Coatomer subunit delta OS=Rattus norvegicus GN=Arcn1 PE=2 SV=1
202	sp Q11011 PSA_MOUSE	Puromycin-sensitive aminopeptidase OS=Mus musculus GN=Npepps PE=1 SV=2
203	sp P11679 K2C8_MOUSE	Keratin, type II cytoskeletal 8 OS=Mus musculus GN=Krt8 PE=1 SV=4
204	sp P02639 S10A1_BOVIN	Protein S100-A1 OS=Bos taurus GN=S100A1 PE=1 SV=2
205	sp Q29443 TRFE_BOVIN	Serotransferrin OS=Bos taurus GN=TF PE=2 SV=1
206	sp Q8VHK6 FLNC_MOUSE	Filamin-C OS=Mus musculus GN=Flnc PE=1 SV=3
207	sp Q76M96 CCD80_HUMAN	Coiled-coil domain-containing protein 80 OS=Homo sapiens GN=CCDC80 PE=1 SV=1
208	sp Q9POL0 VAPA_HUMAN	Vesicle-associated membrane protein-associated protein A OS=Homo sapiens GN=VAPA PE=1 SV=3
209	sp P13355 GTR1_RABIT	Solute carrier family 2, facilitated glucose transporter member 1 OS=Oryctolagus cuniculus GN=SLC2A1 PE=2 SV=1
210	sp Q62159 RHO_C_MOUSE	Rho-related GTP-binding protein RhoC OS=Mus musculus GN=Rhoc PE=1 SV=2
211	sp Q08529 CAN2_MOUSE	Calpain-2 catalytic subunit OS=Mus musculus GN=Capn2 PE=2 SV=4
212	sp Q9TQX6 ACBP_CANFA	Acyl-CoA-binding protein OS=Canis familiaris GN=DBI PE=1 SV=2
213	sp P40227 TCPZ_HUMAN	T-complex protein 1 subunit zeta OS=Homo sapiens GN=CTC6A PE=1 SV=3
214	sp Q5RBU8 ROA2_PONAB	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Pongo abelii GN=HNRNPA2B1 PE=2 SV=1
215	sp Q8BG05 ROA3_MOUSE	Heterogeneous nuclear ribonucleoprotein A3 OS=Mus musculus GN=Hnrnpa3 PE=1 SV=1
216	sp Q5RER0 ITIHS_PONAB	Inter-alpha-trypsin inhibitor heavy chain H5 OS=Pongo abelii GN=ITIHS PE=2 SV=1
217	sp Q8R491 EHD3_RAT	EH domain-containing protein 3 OS=Rattus norvegicus GN=Ehd3 PE=1 SV=2
218	sp Q00612 G6PD1_MOUSE	Glucose-6-phosphate 1-dehydrogenase X OS=Mus musculus GN=G6pdx PE=1 SV=3
219	sp P79136 CAP2B_BOVIN	F-actin-capping protein subunit beta OS=Bos taurus GN=CAP2B PE=1 SV=1
220	sp Q76M23 ZAAA_MOUSE	Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha isoform OS=Mus musculus GN=Ppp2r1a PE=1 SV=3
221	sp P06151 LDHA_MOUSE	L-lactate dehydrogenase A chain OS=Mus musculus GN=Ldha PE=1 SV=3
222	sp Q9GKX8 H590B_HORSE	Heat shock protein HSP 90-beta OS=Equus caballus GN=HSP90AB1 PE=2 SV=3
223	sp Q6SQH4 S10AA_RABIT	Protein S100-A10 OS=Oryctolagus cuniculus GN=S100A10 PE=3 SV=3
224	sp Q76I81 RS12_BOVIN	40S ribosomal protein S12 OS=Bos taurus GN=RP512 PE=2 SV=1
225	sp A7MB70 SEM3C_BOVIN	Semaphorin-3C OS=Bos taurus GN=SEMA3C PE=2 SV=1
226	sp Q9Z1N5 UAP56_MOUSE	Spliceosome RNA helicase Bat1 OS=Mus musculus GN=Bat1 PE=1 SV=1
227	sp P82460 THIO_PIG	Thioredoxin OS=Sus scrofa GN=TXN PE=1 SV=3
228	sp Q92743 HTRA1_HUMAN	Serine protease HTRA1 OS=Homo sapiens GN=HTRA1 PE=1 SV=1
229	sp P04937-2 FINC_RAT	Isoform FNIII-13-less of Fibronectin OS=Rattus norvegicus GN=Fn1
230	sp P09488 GSTM1_HUMAN	Glutathione S-transferase Mu 1 OS=Homo sapiens GN=GSTM1 PE=1 SV=3
231	sp P07737 PROF1_HUMAN	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2
232	sp Q4R502 IDHP_MACFA	Isocitrate dehydrogenase [NADP], mitochondrial OS=Macaca fascicularis GN=IDH2 PE=2 SV=1
233	sp Q27972 CHAD_BOVIN	Chondroadherin OS=Bos taurus GN=CHAD PE=1 SV=1
234	sp Q5R5F0 LDHA_PONAB	L-lactate dehydrogenase A chain OS=Pongo abelii GN=LDHA PE=2 SV=3
235	sp P19001 K1C19_MOUSE	Keratin, type I cytoskeletal 19 OS=Mus musculus GN=Krt19 PE=2 SV=1
236	sp P16403 H12_HUMAN	Histone H1.2 OS=Homo sapiens GN=HIST1H1C PE=1 SV=2
237	sp P45591 COF2_MOUSE	Cofilin-2 OS=Mus musculus GN=Cf2 PE=1 SV=1
238	sp Q8CIB5 FERM2_MOUSE	Fermitin family homolog 2 OS=Mus musculus GN=Fermt2 PE=1 SV=1
239	sp Q9Z2K9 IDHC_MICME	Isocitrate dehydrogenase [NADP] cytoplasmic OS=Microtus mexicanus GN=IDH1 PE=2 SV=1
240	sp Q5E9E6 RL10A_BOVIN	60S ribosomal protein L10a OS=Bos taurus GN=RPL10A PE=2 SV=3
241	sp P12830 CADH1_HUMAN	Cadherin-1 OS=Homo sapiens GN=CDH1 PE=1 SV=3
242	sp P28075 PSB5_RAT	Proteasome subunit beta type-5 OS=Rattus norvegicus GN=Psmb5 PE=1 SV=3
243	sp Q8CFN2 CDC42_RAT	Cell division control protein 42 homolog OS=Rattus norvegicus GN=Cdc42 PE=1 SV=2
244	sp P08294 SODE_HUMAN	Extracellular superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD3 PE=1 SV=2
245	sp Q3ZBD7 G6PI_BOVIN	Glucose-6-phosphate isomerase OS=Bos taurus GN=GPI PE=2 SV=3
246	sp Q14816 ARPC4_BOVIN	Actin-related protein 2/3 complex subunit 4 OS=Bos taurus GN=ARPC4 PE=1 SV=3
247	sp O00300 TR11B_HUMAN	Tumor necrosis factor receptor superfamily member 11B OS=Homo sapiens GN=TNFRSF11B PE=1 SV=2
248	sp P50503 F10A1_RAT	Hsc70-interacting protein OS=Rattus norvegicus GN=St13 PE=1 SV=1
249	sp Q5R514 UBE2L3_PONAB	Ubiquitin-conjugating enzyme E2 L3 OS=Pongo abelii GN=UBE2L3 PE=2 SV=1
250	sp Q9D0M5 DYL2_MOUSE	Dynein light chain 2, cytoplasmic OS=Mus musculus GN=Dynll2 PE=1 SV=1
251	sp P21741 MK_HUMAN	Midkine OS=Homo sapiens GN=MDK PE=1 SV=1

252	sp P50518 VATE1_MOUSE	V-type proton ATPase subunit E 1 OS=Mus musculus GN=Atp6v1e1 PE=1 SV=2
253	sp Q9TTY1 TIMP2_CANFA	Metalloproteinase inhibitor 2 OS=Canis familiaris GN=TIMP2 PE=2 SV=2
254	sp Q71LE2 H33_PIG	Histone H3.3 OS=Sus scrofa GN=H3F3A PE=2 SV=3
255	sp P55918 MFAP4_BOVIN	Microfibril-associated glycoprotein 4 OS=Bos taurus GN=MFAP4 PE=1 SV=2
256	sp Q5RK11 IF4A2_RAT	Eukaryotic initiation factor 4A-II OS=Rattus norvegicus GN=EIF4a2 PE=1 SV=1
257	sp Q86Y38 XYLT1_HUMAN	Xylosyltransferase 1 OS=Homo sapiens GN=XYLT1 PE=1 SV=1
258	sp Q15149-4 PLEC_HUMAN	Isoform Plectin-11 of Plectin OS=Homo sapiens GN=PLEC
259	sp Q6EWQ7 IF5A1_BOVIN	Eukaryotic translation initiation factor 5A-1 OS=Bos taurus GN=EIF5A PE=2 SV=3
260	sp Q9TV56 S10A4_CANFA	Protein S100-A4 OS=Canis familiaris GN=S100A4 PE=3 SV=1
261	sp Q29116 TENA_PIG	Tenascin OS=Sus scrofa GN=TNC PE=1 SV=1
262	sp Q865V6 CAPG_BOVIN	Macrophage-capping protein OS=Bos taurus GN=CAPG PE=2 SV=1
263	sp Q75083 WDR1_HUMAN	WD repeat-containing protein 1 OS=Homo sapiens GN=WDR1 PE=1 SV=4
264	sp Q2EN75 S10A6_PIG	Protein S100-A6 OS=Sus scrofa GN=S100A6 PE=3 SV=1
265	sp Q60847 COCA1_MOUSE	Collagen alpha-1(XII) chain OS=Mus musculus GN=Col12a1 PE=2 SV=3
266	sp Q6UVK1 CSPG4_HUMAN	Chondroitin sulfate proteoglycan 4 OS=Homo sapiens GN=CSPG4 PE=1 SV=2
267	sp O35987 NSF1C_RAT	NSF1 cofactor p47 OS=Rattus norvegicus GN=Nsf1c PE=1 SV=1
268	sp Q5R556 RAN_PONAB	GTP-binding nuclear protein Ran OS=Pongo abelii GN=RAN PE=2 SV=3
269	sp P31324 KAP3_MOUSE	cAMP-dependent protein kinase type II-beta regulatory subunit OS=Mus musculus GN=Prkar2b PE=1 SV=3
270	sp Q9H361 PABP3_HUMAN	Polyadenylate-binding protein 3 OS=Homo sapiens GN=PABPC3 PE=1 SV=2
271	sp O11780 BGH3_PIG	Transforming growth factor-beta-induced protein ig-h3 OS=Sus scrofa GN=TGFBI PE=1 SV=1
272	sp Q07954 LRP1_HUMAN	Prolow-density lipoprotein receptor-related protein 1 OS=Homo sapiens GN=LRP1 PE=1 SV=1
273	sp Q96J81 DYH8_HUMAN	Dynein heavy chain 8, axonemal OS=Homo sapiens GN=DNAH8 PE=1 SV=2
274	sp Q3ZC84 CNDP2_BOVIN	Cytosolic non-specific dipeptidase OS=Bos taurus GN=CNDP2 PE=2 SV=1
275	sp P51635 AK1A1_RAT	Alcohol dehydrogenase [NADP+] OS=Rattus norvegicus GN=Akr1a1 PE=1 SV=2
276	sp P97927 LAMA4_MOUSE	Laminin subunit alpha-4 OS=Mus musculus GN=Lama4 PE=1 SV=2
277	sp Q13822-2 ENPP2_HUMAN	Isoform Alpha of Ectonucleotide pyrophosphatase/phosphodiesterase family member 2 OS=Homo sapiens GN=ENPP2
278	sp Q03001-9 BPA1_HUMAN	Isoform EB of Bullous pemphigoid antigen 1 OS=Homo sapiens GN=DST
279	sp P11586 C1TC_HUMAN	C-1-tetrahydrofolate synthase, cytoplasmic OS=Homo sapiens GN=MTHFD1 PE=1 SV=3
280	sp P14866 HNRPL_HUMAN	Heterogeneous nuclear ribonucleoprotein L OS=Homo sapiens GN=HNRNPL PE=1 SV=2
281	sp Q9NRR5 OLF3_HUMAN	Olfactomedin-like protein 3 OS=Homo sapiens GN=OLFML3 PE=2 SV=1
282	sp Q4V8H8 EHD2_RAT	EH domain-containing protein 2 OS=Rattus norvegicus GN=Ehd2 PE=1 SV=1
283	sp Q7TMV8 HUWE1_MOUSE	E3 ubiquitin-protein ligase HUWE1 OS=Mus musculus GN=Huwe1 PE=1 SV=4
284	sp P00570 KAD1_BOVIN	Adenylate kinase isoenzyme 1 OS=Bos taurus GN=AK1 PE=1 SV=2
285	sp Q6P502 TCPG_RAT	T-complex protein 1 subunit gamma OS=Rattus norvegicus GN=Cct3 PE=1 SV=1
286	sp P00813 ADA_HUMAN	Adenosine deaminase OS=Homo sapiens GN=ADA PE=1 SV=3
287	sp Q3T035 ARPC3_BOVIN	Actin-related protein 2/3 complex subunit 3 OS=Bos taurus GN=ARPC3 PE=1 SV=3
288	sp P46940 IQGA1_HUMAN	Ras GTPase-activating-like protein IQGAP1 OS=Homo sapiens GN=IQGAP1 PE=1 SV=1
289	sp Q9TSZ6 DAG1_CANFA	Dystroglycan OS=Canis familiaris GN=DAG1 PE=3 SV=1

290	sp Q5RBG1 RAB5B_PONAB	Ras-related protein Rab-5B OS=Pongo abelii GN=RAB5B PE=2 SV=1
291	sp P70349 HINT1_MOUSE	Histidine triad nucleotide-binding protein 1 OS=Mus musculus GN=Hint1 PE=1 SV=3
292	sp Q08D0P PGM1_BOVIN	Phosphoglucosyltransferase-1 OS=Bos taurus GN=PGM1 PE=2 SV=1
293	sp Q9TUL9 TIMP3_HORSE	Metalloproteinase inhibitor 3 OS=Equus caballus GN=TIMP3 PE=2 SV=1
294	sp P49747 COMP_HUMAN	Cartilage oligomeric matrix protein OS=Homo sapiens GN=COMP PE=1 SV=2
295	sp Q5VYK3 ECM29_HUMAN	Proteasome-associated protein ECM29 homolog OS=Homo sapiens GN=ECM29 PE=1 SV=2
296	sp Q63210 GNA12_RAT	Guanine nucleotide-binding protein subunit alpha-12 OS=Rattus norvegicus GN=Gna12 PE=2 SV=3
297	sp P15170 ERF3A_HUMAN	Eukaryotic peptide chain release factor GTP-binding subunit ERF3A OS=Homo sapiens GN=GSPT1 PE=1 SV=1
298	sp P52174 NDKA1_BOVIN	Nucleoside diphosphate kinase A 1 OS=Bos taurus GN=NME1-1 PE=1 SV=3
299	sp Q4VIT4 PDIA3_CERAE	Protein disulfide-isomerase A3 OS=Cercopithecus aethiops GN=PDIA3 PE=2 SV=1
300	sp Q6AY84 SCRN1_RAT	Secernin-1 OS=Rattus norvegicus GN=Scrn1 PE=1 SV=1
301	sp Q5E984 TCTP_BOVIN	Translationally-controlled tumor protein OS=Bos taurus GN=TPT1 PE=2 SV=1
302	sp P15311 EZRI_HUMAN	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4
303	sp Q9JHW0 PSB7_RAT	Proteasome subunit beta type-7 OS=Rattus norvegicus GN=Psb7 PE=1 SV=1
304	sp Q9UL16 CCD19_HUMAN	Coiled-coil domain-containing protein 19, mitochondrial OS=Homo sapiens GN=CCDC19 PE=1 SV=2
305	sp Q62920 PDLI5_RAT	PDZ and LIM domain protein 5 OS=Rattus norvegicus GN=Pdlm5 PE=1 SV=2
306	sp P30044 PRDX5_HUMAN	Peroxisiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5 PE=1 SV=3
307	sp Q9HBL0 TENS1_HUMAN	Tensin-1 OS=Homo sapiens GN=TN51 PE=1 SV=2
308	sp Q02809 PLOD1_HUMAN	Procollagen-lysine,2-oxoglutarate 5-dioxygenase 1 OS=Homo sapiens GN=PLOD1 PE=1 SV=2
309	sp P63322 RALA_RAT	Ras-related protein Ral-A OS=Rattus norvegicus GN=Rala PE=1 SV=1
310	sp Q5RB59 PUR6_PONAB	Multifunctional protein ADE2 OS=Pongo abelii GN=PAICS PE=2 SV=3
311	sp Q86V81 THOC4_HUMAN	THO complex subunit 4 OS=Homo sapiens GN=THOC4 PE=1 SV=3
312	sp Q9CPW4 ARPC5_MOUSE	Actin-related protein 2/3 complex subunit 5 OS=Mus musculus GN=Arpc5 PE=2 SV=3
313	sp Q88I20 PCD20_MOUSE	Protocadherin-20 OS=Mus musculus GN=Pcdh20 PE=2 SV=2
314	sp P20072 ANXA7_BOVIN	Annexin A7 OS=Bos taurus GN=ANXA7 PE=1 SV=2
315	sp Q9H223 EHD4_BOVIN	EH domain-containing protein 4 OS=Homo sapiens GN=EHD4 PE=1 SV=1
316	sp Q8WNW3 PLAK_PIG	Junction plakoglobin OS=Sus scrofa GN=Jup PE=2 SV=1
317	sp Q9IK81 MYG1_MOUSE	UPF0160 protein MYG1, mitochondrial OS=Mus musculus GN=Myg1 PE=2 SV=1
318	sp P17980 PR56A_HUMAN	26S protease regulatory subunit 6A OS=Homo sapiens GN=PSMC3 PE=1 SV=3
319	sp Q5EAC6 CDC37_BOVIN	Hsp90 co-chaperone Cdc37 OS=Bos taurus GN=CDC37 PE=2 SV=1
320	sp Q9CV86 ARPC2_MOUSE	Actin-related protein 2/3 complex subunit 2 OS=Mus musculus GN=Arpc2 PE=1 SV=3
321	sp P50995 ANX11_HUMAN	Annexin A11 OS=Homo sapiens GN=ANXA11 PE=1 SV=1
322	sp Q95204 CY561_SHEEP	Cytochrome b561 OS=Ovis aries GN=Cyb561 PE=2 SV=1
323	sp P79342 S10AD_BOVIN	Protein S100-A13 OS=Bos taurus GN=S100A13 PE=3 SV=2
324	sp Q5NV72 ZN770_PONAB	Zinc finger protein 770 (Fragment) OS=Pongo abelii GN=ZNF770 PE=2 SV=1
325	sp Q4PLI0 NEDD8_RABIT	NEDD8 OS=Oryctolagus cuniculus GN=NEDD8 PE=3 SV=1
326	sp Q920N1 IF2G_MOUSE	Eukaryotic translation initiation factor 2 subunit 3, X-linked OS=Mus musculus GN=EIF2S3X PE=1 SV=2
327	sp Q99969 RARR2_HUMAN	Retinoic acid receptor responder protein 2 OS=Homo sapiens GN=RARRS2 PE=1 SV=1
328	sp Q6YKA4 HMGB1_CANFA	High mobility group protein B1 OS=Canis familiaris GN=HMGB1 PE=2 SV=3
329	sp Q9N0Y0 VAMP2_MACMU	Vesicle-associated membrane protein 2 OS=Macaca mulatta GN=VAMP2 PE=3 SV=3
330	sp Q16881 TXNR1_HUMAN	Thioredoxin reductase 1, cytoplasmic OS=Homo sapiens GN=TXNRD1 PE=1 SV=3
331	sp Q32C42 ADHX_BOVIN	Alcohol dehydrogenase class-3 OS=Bos taurus GN=ADH5 PE=2 SV=1
332	sp P33097 AATC_BOVIN	Aspartate aminotransferase, cytoplasmic OS=Bos taurus GN=GOT1 PE=1 SV=3
333	RRRRRsp AQCPA3 BTBD1_BOVIN	REVERSED BTB/POZ domain-containing protein 19 OS=Bos taurus GN=BTBD19 PE=2 SV=1
334	sp P27816 MAP4_HUMAN	Microtubule-associated protein 4 OS=Homo sapiens GN=MAP4 PE=1 SV=3
335	sp Q5R5U1 RAB10_PONAB	Ras-related protein Rab-10 OS=Pongo abelii GN=RAB10 PE=2 SV=1
336	sp Q68FQ0 TCPE_RAT	T-complex protein 1 subunit epsilon OS=Rattus norvegicus GN=Cct5 PE=1 SV=1
337	sp Q5RAE1 CPNE3_PONAB	Copine-3 OS=Pongo abelii GN=CPNE3 PE=2 SV=1
338	RRRRRsp Q9D700 DUS26_MOUSE	REVERSED Dual specificity protein phosphatase 26 OS=Mus musculus GN=Dusp26 PE=2 SV=2
339	sp Q75339 CILP1_HUMAN	Cartilage intermediate layer protein 1 OS=Homo sapiens GN=CILP PE=1 SV=3
340	sp Q9UBG0 MRC2_HUMAN	C-type mannose receptor 2 OS=Homo sapiens GN=MRC2 PE=1 SV=1
341	sp Q60763 USO1_HUMAN	General vesicular transport factor p115 OS=Homo sapiens GN=USO1 PE=1 SV=2
342	sp P16086 SPTA2_RAT	Spectrin alpha chain, brain OS=Rattus norvegicus GN=Sptan1 PE=1 SV=2
343	sp Q5BJY9 K1C18_RAT	Keratin, type I cytoskeletal 18 OS=Rattus norvegicus GN=Krt18 PE=1 SV=3
344	sp P62909 RS3_RAT	40S ribosomal protein S3 OS=Rattus norvegicus GN=Rps3 PE=2 SV=1
345	sp P80313 TCPH_MOUSE	T-complex protein 1 subunit eta OS=Mus musculus GN=Cct7 PE=1 SV=1
346	sp Q95117 FRP3_BOVIN	Secreted frizzled-related protein 3 OS=Bos taurus GN=FRZB PE=1 SV=1
347	sp Q641Y8 DDX1_RAT	ATP-dependent RNA helicase DDX1 OS=Rattus norvegicus GN=Ddx1 PE=2 SV=1
348	RRRRRsp Q9H255 RNF39_HUMAN	REVERSED RING finger protein 39 OS=Homo sapiens GN=RNF39 PE=1 SV=2
349	sp Q9R1P1 PSB3_MOUSE	Proteasome subunit beta type-3 OS=Mus musculus GN=Psb3 PE=1 SV=1
350	sp Q3SZF2 ARF4_BOVIN	ADP-ribosylation factor 4 OS=Bos taurus GN=ARF4 PE=2 SV=3
351	sp P05064 ALDOA_MOUSE	Fructose-bisphosphate aldolase A OS=Mus musculus GN=Aldoa PE=1 SV=2
352	sp Q04857 C06A1_MOUSE	Collagen alpha-1(VI) chain OS=Mus musculus GN=Col6a1 PE=2 SV=1
353	sp Q4R572 1433B_MACFA	14-3-3 protein beta/alpha OS=Macaca fascicularis GN=YWHAB PE=2 SV=3
354	sp Q68FP1 GELS_RAT	Gelsolin OS=Rattus norvegicus GN=Gsn PE=1 SV=1
355	sp Q8TER5-3 SOLO_HUMAN	Isoform 3 of Protein SOLO OS=Homo sapiens GN=SOLO
356	sp P35221-2 CTNA1_HUMAN	Isoform 2 of Catenin alpha-1 OS=Homo sapiens GN=CTNNA1
357	sp Q10994 CYTB_SHEEP	Cystatin-B OS=Ovis aries GN=CSTB PE=1 SV=1
358	sp Q43242 PSMD3_HUMAN	26S proteasome non-ATPase regulatory subunit 3 OS=Homo sapiens GN=PSMD3 PE=1 SV=2
359	sp Q01634 IDUA_CANFA	Alpha-L-iduronidase OS=Canis familiaris GN=IDUA PE=1 SV=1
360	sp Q60692 PSB6_MOUSE	Proteasome subunit beta type-6 OS=Mus musculus GN=Psb6 PE=1 SV=3
361	sp P78417 GSTO1_HUMAN	Glutathione S-transferase omega-1 OS=Homo sapiens GN=GSTO1 PE=1 SV=2
362	sp P47959 HPRT_MERUN	Hypoxanthine-guanine phosphoribosyltransferase OS=Meriones unguiculatus GN=HPRT1 PE=2 SV=2
363	sp Q5T7N8 F27D1_HUMAN	Protein FAM27D1 OS=Homo sapiens GN=FAM27D1 PE=3 SV=2
364	sp Q9XSJ4 ENOA_BOVIN	Alpha-enolase OS=Bos taurus GN=ENO1 PE=1 SV=4
365	sp Q9H4M9 EHD1_HUMAN	EH domain-containing protein 1 OS=Homo sapiens GN=EHD1 PE=1 SV=2
366	sp P07450 CAH3_HORSE	Carbonic anhydrase 3 OS=Equus caballus GN=CA3 PE=1 SV=2
367	sp Q63610 TPM3_RAT	Tropomyosin alpha-3 chain OS=Rattus norvegicus GN=Tpm3 PE=1 SV=2
368	sp Q9NQ57-2 INCE_HUMAN	Isoform 2 of Inner centromere protein OS=Homo sapiens GN=INCENP
369	sp Q70435 PSA3_MOUSE	Proteasome subunit alpha type-3 OS=Mus musculus GN=Psa3 PE=1 SV=3
370	sp Q70492 SNX3_MOUSE	Sorting nexin-3 OS=Mus musculus GN=Snx3 PE=2 SV=3
371	sp Q9WV84 SLIT3_MOUSE	Slit homolog 3 protein OS=Mus musculus GN=Slit3 PE=2 SV=1
372	sp Q43252 PAPS1_HUMAN	Bifunctional 3'-phosphoadenosine 5'-phosphosulfate synthase 1 OS=Homo sapiens GN=PAPS1 PE=1 SV=2
373	sp Q4R416 CAP1_MACFA	Adenylyl cyclase-associated protein 1 OS=Macaca fascicularis GN=CAP1 PE=2 SV=3
374	sp P47199 QOR_MOUSE	Quinone oxidoreductase OS=Mus musculus GN=Cryz PE=2 SV=1
375	sp Q61481 PDE1A_MOUSE	Calcium/calmodulin-dependent 3',5'-cyclic nucleotide phosphodiesterase 1A OS=Mus musculus GN=Pde1a PE=2 SV=1
376	sp Q92204 HNRPC_MOUSE	Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Mus musculus GN=Hnrnpc PE=1 SV=1
377	sp Q6AY09 HNRH2_RAT	Heterogeneous nuclear ribonucleoprotein H2 OS=Rattus norvegicus GN=Hnrhp2 PE=1 SV=1