

GeneBank access	<i>p</i> -value ANOVA	Best hit description	Best hit evalue
AJ427420	1,00E-05	(sp O35607) Bone morphogenetic protein receptor type-2 OS=Mus musculus GN=Bmpr2	9,00E-97
AJ544884	1,00E-05	(sp Q9R229) Bone morphogenetic protein 10 OS=Mus musculus GN=Bmp10 (sp Q0P4Y4) Calcitonin gene-related peptide type 1 receptor OS=Xenopus tropicalis GN=calcr1	7,00E-44
AJ551182	1,00E-06		1,00E-50
AJ565449	1,00E-06		
AJ565474	1,00E-06	hypothetical protein AaeL_AAEL013387 [Aedes aegypti]	8,00E-06
AJ565476	1,00E-05	(sp P35980) 60S ribosomal protein L18 OS=Mus musculus GN=Rpl18 (sp Q863A5) Dyslexia susceptibility 1 candidate gene 1 protein homolog OS=Gorilla gorilla gorilla GN=DYX1C1	4,00E-70
AJ565518	1,00E-06		7,00E-45
AJ565526	1,00E-06		
AJ565540	1,00E-05		
AJ565553	1,00E-05	(sp Q2NKT1) Profilin-4 OS=Bos taurus GN=PFN4 (sp Q80W96) Iron-sulfur cluster assembly 1 homolog, mitochondrial OS=Rattus norvegicus GN=Isca1	1,00E-06
AJ565559	1,00E-06		9,00E-46
AJ565564	1,00E-06		
AJ565570	1,00E-06	(sp P00399) Cytochrome c oxidase subunit 1 OS=Drosophila melanogaster GN=mt:Col	1,00E-137
AJ565585	1,00E-06		
AJ565599	1,00E-06	(sp Q641Z5) Cation transport regulator-like protein 2 OS=Rattus norvegicus GN=Chac2	3,00E-34
AJ565605	1,00E-05		
AJ565651	1,00E-06	(sp Q53G44) Interferon-induced protein 44-like OS=Homo sapiens GN=IFI44L	7,00E-33
AJ565656	1,00E-06		
AJ565678	1,00E-06	(sp Q16981) Aplycalcin OS=Aplysia californica	3,00E-55
AJ565682	1,00E-05		
AJ565688	1,00E-06	hypothetical protein BBAL3_1222 [Brevundimonas sp. BAL3]	2,00E-26
AJ565701	1,00E-06	Gallus gallus heat shock protein 25 (HSP25), mRNA	8,00E-08
AJ565712	1,00E-06		
AJ565773	1,00E-06		
AJ565812	1,00E-06		

AJ890150	1,00E-05	(sp O42329) Gonadotropin-releasing hormone II receptor OS=Clarias gariepinus	3,00E-37
AM237711	1,00E-06		
AM237717	1,00E-06		
AM237736	1,00E-05		
AM237745	1,00E-06		
AM237747	1,00E-06		
AM237756	1,00E-06		
AM237757	1,00E-06		
AM237765	1,00E-06		
AM237777	1,00E-05		
AM237784	1,00E-05		
AM237786	1,00E-06		
AM853227	1,00E-06	(sp P02840) Salivary glue protein Sgs-3 OS=Drosophila melanogaster GN=Sgs3	2,00E-11
AM853250	1,00E-06		
AM853258	1,00E-05	(sp P52746) Zinc finger protein 142 OS=Homo sapiens GN=ZNF142	7,00E-24
AM853261	1,00E-06	(sp Q4KLU2) Pre-mRNA-processing factor 39 OS=Xenopus laevis GN=prpf39	1,00E-68
AM853272	1,00E-06		
AM853277	1,00E-06	(sp Q7Z5L9) Interferon regulatory factor 2-binding protein 2 OS=Homo sapiens GN=IRF2BP2	6,00E-39
AM853285	1,00E-06		
AM853310	1,00E-06	(sp Q08420) Extracellular superoxide dismutase [Cu-Zn] OS=Rattus norvegicus GN=Sod3	8,00E-10
AM853313	1,00E-05		
AM853315	1,00E-06		
AM853317	1,00E-06		
AM853325	1,00E-06		
AM853327	1,00E-06	(sp Q9WVB4) Slit homolog 3 protein OS=Mus musculus GN=Slit3	4,00E-09
AM853335	1,00E-05		
AM853347	1,00E-06		
AM853353	1,00E-06	(sp Q5RFB0) Aminoacylase-1 OS=Pongo abelii GN=ACY1	3,00E-67
AM853357	1,00E-05		
AM853376	1,00E-06	(sp Q7YRL2) Calponin-1 OS=Ovis aries GN=CNN1	9,00E-24

AM853378	1,00E-06		
AM853383	1,00E-06	(sp Q8NA69) Uncharacterized protein C19orf45 OS=Homo sapiens GN=C19orf45	4,00E-07
AM853386	1,00E-06		
AM853390	1,00E-06		
AM853391	1,00E-06		
AM853403	1,00E-06	(sp Q00945) Conopressin/neurophysin OS=Lymnaea stagnalis (sp Q81ZJ3) C3 and PZP-like alpha-2-macroglobulin domain-containing protein 8 OS=Homo sapiens GN=CPAMD8	1,00E-32
AM853418	1,00E-05		
AM853432	1,00E-06		
AM853434	1,00E-05		
AM853438	1,00E-06		
AM853488	1,00E-06	(sp Q99MK2) Histone acetyltransferase KAT5 OS=Rattus norvegicus GN=Kat5	4,00E-17
AM853500	1,00E-06		
AM853505	1,00E-06		
AM853518	1,00E-06	(sp Q5ZK69) Proteasomal ATPase-associated factor 1 OS=Gallus gallus GN=PAAF1	6,00E-54
AM853528	1,00E-06		
AM853530	1,00E-06		
AM853544	1,00E-06		
AM853546	1,00E-06		
AM853548	1,00E-06		
AM853593	1,00E-06		
AM853601	1,00E-06		
AM853602	1,00E-06		
AM853613	1,00E-06	(sp Q5VYJ5) MAM and LDL-receptor class A domain-containing protein C10orf112 OS=Homo sapiens	1,00E-36
AM853618	1,00E-06	(sp P04040) Catalase OS=Homo sapiens GN=CAT	1,00E-101
AM853624	1,00E-06		
AM853626	1,00E-06	(sp O75911) Short-chain dehydrogenase/reductase 3 OS=Homo sapiens GN=DHRS3	1,00E-46
AM853633	1,00E-06		
AM853637	1,00E-06	(sp Q8VHC8) Lysosomal alpha-mannosidase OS=Cavia porcellus GN=MAN2B1	4,00E-39

AM853638	1,00E-06		
AM853643	1,00E-06	(sp Q94887) Neurexin-4 OS=Drosophila melanogaster GN=Nrx-IV	1,00E-08
AM853645	1,00E-06		
AM853655	1,00E-06	(sp P09057) Ornithine decarboxylase OS=Rattus norvegicus GN=Odc1	4,00E-95
AM853662	1,00E-05		
AM853689	1,00E-05		
AM853709	1,00E-06		
AM853723	1,00E-05	(sp Q3MHR7) Actin-related protein 2/3 complex subunit 2 OS=Bos taurus GN=ARPC2	1,00E-132
AM853724	1,00E-06	(sp Q28FH2) N-terminal kinase-like protein OS=Xenopus tropicalis GN=scyl1	3,00E-18
AM853725	1,00E-05	(sp O19112) Cartilage intermediate layer protein 1 (Fragment) OS=Sus scrofa GN=CILP	2,00E-09
AM853754	1,00E-05		
AM853767	1,00E-06		
AM853782	1,00E-06		
AM853786	1,00E-05	(sp Q1PRL4) Tripartite motif-containing protein 71 OS=Gallus gallus GN=TRIM71	1,00E-08
AM853794	1,00E-06	(sp P46684) Zinc finger protein ZIC 1 OS=Mus musculus GN=Zic1	1,00E-22
AM853802	1,00E-06	hypothetical protein IscW_ISCW013676 [Ixodes scapularis]	9,00E-13
AM853803	1,00E-06		
AM853818	1,00E-06		
AM853829	1,00E-06		
AM853830	1,00E-06		
AM853831	1,00E-05		
AM853833	1,00E-06		
AM853835	1,00E-06		
AM853838	1,00E-06	PREDICTED: similar to lethal (2) k09913 CG3082-PC [Danio rerio]	2,00E-28
AM853851	1,00E-06		
AM853859	1,00E-05		
AM853862	1,00E-06		
AM853867	1,00E-06		
AM853869	1,00E-06	(sp Q0P4F7) Acyl-CoA synthetase family member 2, mitochondrial OS=Danio rerio GN=acsf2	4,00E-06
AM853870	1,00E-06		

AM853880	1,00E-06		
AM853888	1,00E-06		
AM853905	1,00E-05	(sp Q4R7L8) Peroxisomal NADH pyrophosphatase NUDT12 OS=Macaca fascicularis GN=NUDT12	3,00E-45
AM853912	1,00E-05		
AM853929	1,00E-06	(sp Q95044) Ubiquitin-conjugating enzyme E2 C OS=Spisula solidissima	3,00E-62
AM853931	1,00E-06	(sp P49165) 60S ribosomal protein L4 OS=Urechis caupo GN=RPL4	1,00E-159
AM853938	1,00E-06	(sp Q6ZQY2) Leucine-rich repeat-containing protein LOC400891 OS=Homo sapiens	7,00E-23
AM853940	1,00E-06	(sp O18404) 3-hydroxyacyl-CoA dehydrogenase type-2 OS=Drosophila melanogaster GN=scu	2,00E-36
AM853942	1,00E-06		
AM853949	1,00E-06		
AM853951	1,00E-05		
AM853954	1,00E-05		
AM853956	1,00E-06		
AM853957	1,00E-06	C. briggsae CBR-LNTL-1 protein [Caenorhabditis briggsae]	6,00E-27
AM853970	1,00E-06		
AM853974	1,00E-05	(sp Q9BYK8) Peroxisomal proliferator-activated receptor A-interacting complex 285 kDa protein OS=Homo sapiens	9,00E-64
AM853975	1,00E-05	(sp O35125) Leucine-rich repeat-containing protein 23 OS=Mus musculus GN=Lrrc23	7,00E-62
AM854003	1,00E-06	(sp P34528) Putative serine protease K12H4.7 OS=Caenorhabditis elegans GN=K12H4.7	7,00E-32
AM854028	1,00E-06	(sp P13608) Aggrecan core protein OS=Bos taurus GN=ACAN	3,00E-12
AM854029	1,00E-06	(sp P52161) Protein max OS=Danio rerio GN=max	4,00E-36
AM854037	1,00E-06	(sp P06144) Late histone H1 OS=Lytechinus pictus	2,00E-18
AM854060	1,00E-06		
AM854063	1,00E-05		
AM854064	1,00E-06	(sp Q7KRY7) Protein lap4 OS=Drosophila melanogaster GN=scrib	4,00E-07
AM854067	1,00E-05		
AM854072	1,00E-06		
AM854078	1,00E-06	(sp Q62288) Testican-1 OS=Mus musculus GN=Spock1	6,00E-30
AM854090	1,00E-06	(sp Q9JL99) C-type lectin domain family 1 member B OS=Mus musculus GN=Clec1b	2,00E-07

AM854097	1,00E-06		
AM854099	1,00E-06	(sp P06603) Tubulin alpha-1 chain OS=Drosophila melanogaster GN=alphaTub84B	1,00E-173
AM854102	1,00E-06	(sp O76756) 40S ribosomal protein S8 OS=Apis mellifera GN=RpS8	2,00E-80
AM854108	1,00E-05	(sp Q99L04) Dehydrogenase/reductase SDR family member 1 OS=Mus musculus GN=Dhrs1	4,00E-82
AM854110	1,00E-06		
AM854117	1,00E-06		
AM854121	1,00E-06		
AM854125	1,00E-06	(sp Q32Q06) AP-1 complex subunit mu-1 OS=Rattus norvegicus GN=Ap1m1	2,00E-17
AM854138	1,00E-06		
AM854157	1,00E-06		
AM854168	1,00E-06		
AM854191	1,00E-06	(sp Q63396) Activated RNA polymerase II transcriptional coactivator p15 OS=Rattus norvegicus GN=Sub1	1,00E-15
AM854192	1,00E-05	(sp Q0D2G3) Transmembrane protein C10orf57 homolog OS=Bos taurus	9,00E-10
AM854224	1,00E-06		
AM854226	1,00E-05		
AM854238	1,00E-06		
AM854259	1,00E-06		
AM854260	1,00E-05		
AM854291	1,00E-06	(sp P62250) 40S ribosomal protein S16 OS=Rattus norvegicus GN=Rps16	1,00E-63
AM854308	1,00E-06		
AM854313	1,00E-06		
AM854316	1,00E-06		
AM854317	1,00E-06	(sp P51688) N-sulphoglucosamine sulphohydrolase OS=Homo sapiens GN=SGSH	3,00E-37
AM854330	1,00E-06		
AM854358	1,00E-06	(sp B3GNI6) Septin-11 OS=Rattus norvegicus GN=Sept11	6,00E-83
AM854365	1,00E-06		
AM854376	1,00E-06	(sp Q24439) ATP synthase subunit O, mitochondrial OS=Drosophila melanogaster GN=Oscp	6,00E-39
AM854380	1,00E-06	(sp Q13263) Transcription intermediary factor 1-beta OS=Homo sapiens GN=TRIM28 SV=5	1,00E-08

AM854386	1,00E-06	(sp Q9BRT3) Uncharacterized protein C17orf37 OS=Homo sapiens GN=C17orf37	1,00E-07
AM854397	1,00E-06		
AM854402	1,00E-06	(sp Q15751) Probable E3 ubiquitin-protein ligase HERC1 OS=Homo sapiens GN=HERC1	2,00E-82
AM854428	1,00E-05	(sp P62870) Transcription elongation factor B polypeptide 2 OS=Rattus norvegicus GN=Tceb2	3,00E-36
AM854439	1,00E-05	(sp P07898) Aggrecan core protein OS=Gallus gallus GN=ACAN	1,00E-21
AM854464	1,00E-05		
AM854472	1,00E-06		
AM854476	1,00E-06		
AM854508	1,00E-06		
AM854528	1,00E-06		
AM854565	1,00E-06	(sp Q6R8J2) Protein HEG OS=Danio rerio GN=heg	3,00E-07
AM854574	1,00E-06		
AM854589	1,00E-06	(sp Q8BGN3) Ectonucleotide pyrophosphatase/phosphodiesterase family member 6 OS=Mus musculus GN=Enpp6	8,00E-06
AM854601	1,00E-06		
AM854604	1,00E-06	(sp Q08013) Translocon-associated protein subunit gamma OS=Rattus norvegicus GN=Ssr3	2,00E-69
AM854608	1,00E-06		
AM854612	1,00E-06		
AM854624	1,00E-05		
AM854635	1,00E-06		
AM854643	1,00E-06		
AM854651	1,00E-06	(sp Q9PTW9) Proteasome subunit alpha type-7 OS=Carassius auratus GN=psma7	1,00E-102
AM854668	1,00E-05	(sp Q28024) Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-12 OS=Bos taurus GN=GNG12	4,00E-08
AM854681	1,00E-06	(sp Q9CRB9) Coiled-coil-helix-coiled-coil-helix domain-containing protein 3, mitochondrial OS=Mus musculus	7,00E-13
AM854684	1,00E-06		
AM854687	1,00E-06		
AM854691	1,00E-06	(sp A5D989) Elongation factor 1-delta OS=Bos taurus GN=EEF1D	1,00E-32
AM854698	1,00E-05		

AM854699	1,00E-06	(sp P58059) 28S ribosomal protein S21, mitochondrial OS=Mus musculus GN=Mrps21	6,00E-12
AM854702	1,00E-06	(sp Q91X83) S-adenosylmethionine synthetase isoform type-1 OS=Mus musculus GN=Mat1a	1,00E-134
AM854704	1,00E-06		
AM854714	1,00E-06	(sp P42577) Soma ferritin OS=Lymnaea stagnalis	3,00E-75
AM854720	1,00E-06	(sp Q60HH4) Acid ceramidase OS=Macaca fascicularis GN=ASAH1	8,00E-85
AM854721	1,00E-06		
AM854723	1,00E-06		
AM854751	1,00E-06	(sp Q5ZIG3) Solute carrier family 25 member 46 OS=Gallus gallus GN=SLC25A46	2,00E-81
AM854755	1,00E-06	(sp Q8R0G9) Nuclear pore complex protein Nup133 OS=Mus musculus GN=Nup133	1,00E-45
AM854772	1,00E-06		
AM854775	1,00E-05	(sp O08997) Copper transport protein ATOX1 OS=Mus musculus GN=Atox1	6,00E-10
AM854779	1,00E-05		
AM854781	1,00E-06	(sp Q9ERZ8) Transient receptor potential cation channel subfamily V member 4 OS=Rattus norvegicus GN=Trpv4	1,00E-06
AM854783	1,00E-06		
AM854792	1,00E-06	(sp Q9IA76) 60S ribosomal protein L31 OS=Paralichthys olivaceus GN=rpl31	8,00E-48
AM854798	1,00E-06	(sp P30337) N-chimaerin OS=Rattus norvegicus GN=Chn1	6,00E-20
AM854799	1,00E-05		
AM854804	1,00E-06		
AM854807	1,00E-05	(sp Q6PHU5) Sortilin OS=Mus musculus GN=Sort1	1,00E-18
AM854812	1,00E-05	(sp A4IH75) Autophagy-related protein 101 OS=Xenopus tropicalis GN=atg101	7,00E-17
AM854820	1,00E-06		
AM854825	1,00E-06		
AM854827	1,00E-06		
AM854835	1,00E-06	(sp Q6PBT7) Mitochondrial import inner membrane translocase subunit TIM14 OS=Danio rerio GN=dnajc19	8,00E-27
AM854836	1,00E-06	PREDICTED: similar to antistasin [Hydra magnipapillata]	6,00E-07
AM854841	1,00E-06	(sp P30050) 60S ribosomal protein L12 OS=Homo sapiens GN=RPL12	2,00E-64

AM854842	1,00E-06	hypothetical protein BRAFLDRAFT_98499 [Branchiostoma floridae]	2,00E-16
AM854843	1,00E-06		
AM854849	1,00E-06	(sp Q5U300) Ubiquitin-like modifier-activating enzyme 1 OS=Rattus norvegicus GN=Uba1	4,00E-54
AM854854	1,00E-06		
AM854856	1,00E-06		
AM854863	1,00E-06		
AM854865	1,00E-06		
AM854869	1,00E-06	(sp Q7M456) Ribonuclease Oy OS=Crassostrea gigas	4,00E-07
AM854871	1,00E-06	(sp Q03105) V-type proton ATPase 16 kDa proteolipid subunit OS=Torpedo marmorata	9,00E-52
AM854880	1,00E-06		
AM854888	1,00E-06	(sp O94581) Cytochrome c oxidase subunit 6B OS=Schizosaccharomyces pombe GN=cox12	2,00E-21
AM854895	1,00E-06		
AM854898	1,00E-06	(sp Q8NFL0) UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 7 OS=Homo sapiens GN=B3GNT7	2,00E-14
AM854900	1,00E-06	(sp A7SMW7) L-2-hydroxyglutarate dehydrogenase, mitochondrial OS=Nematostella vectensis GN=v1g172254	2,00E-58
AM854901	1,00E-06	(sp Q2M2S8) Alkylated DNA repair protein alkB homolog 7 OS=Bos taurus GN=ALKBH7	4,00E-26
AM854905	1,00E-06	(sp Q5U3A7) Dynamin-like 120 kDa protein, mitochondrial OS=Danio rerio GN=opa1	1,00E-56
AM854925	1,00E-06	PREDICTED: hypothetical protein, partial [Acyrtosiphon pisum]	4,00E-09
AM854931	1,00E-06		
AM854934	1,00E-06		
AM854941	1,00E-05	(sp A1L314) Macrophage-expressed gene 1 protein OS=Mus musculus GN=Mpeg1	2,00E-29
AM854958	1,00E-06	(sp Q0VGK4) Glycerophosphodiester phosphodiesterase domain-containing protein 1 OS=Rattus norvegicus	4,00E-30
AM854959	1,00E-06		
AM854974	1,00E-06	(sp Q3TOY3) Sulfotransferase family cytosolic 1B member 1 OS=Bos taurus GN=SULT1B1	5,00E-06
AM855003	1,00E-06		
AM855008	1,00E-06		

AM855015	1,00E-06	(sp P53992) Protein transport protein Sec24C OS=Homo sapiens GN=SEC24C	3,00E-16
AM855019	1,00E-05	hypothetical protein BRAFLDRAFT_76550 [Branchiostoma floridae]	6,00E-25
AM855031	1,00E-06		
		(sp Q8TC84) Fibronectin type 3 and ankyrin repeat domains protein 1	
AM855066	1,00E-05	OS=Homo sapiens GN=FANK1	2,00E-31
AM855080	1,00E-06	(sp Q64012) RNA-binding protein Raly OS=Mus musculus GN=Raly	5,00E-22
AM855089	1,00E-06	(sp Q6E2N3) E3 ubiquitin-protein ligase TRIM33 OS=Danio rerio GN=trim33	7,00E-08
AM855094	1,00E-05		
AM855095	1,00E-06	CG31973, isoform D [Drosophila melanogaster]	4,00E-07
AM855097	1,00E-06	(sp Q3TD49) Signal peptide peptidase-like 2B OS=Mus musculus GN=Sppl2b	3,00E-08
AM855098	1,00E-06		
AM855099	1,00E-05		
		(sp Q6DFM1) SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily B member 1	1,00E-80
AM855106	1,00E-06		
AM855107	1,00E-06		
AM855113	1,00E-06		
AM855114	1,00E-05		
AM855115	1,00E-05	PREDICTED: similar to multiple EGF-like-domains 6 [Ciona intestinalis]	5,00E-06
		(sp B4N9T4) Leucine-rich repeat protein soc-2 homolog OS=Drosophila willistoni GN=Sur-8	3,00E-69
AM855119	1,00E-06		
AM855129	1,00E-06	(sp P02362) 40S ribosomal protein S7 OS=Xenopus laevis GN=rps7	3,00E-39
AM855132	1,00E-06		
AM855144	1,00E-06	PREDICTED: similar to conserved hypothetical protein [Hydra magnipapillata]	3,00E-17
AM855148	1,00E-06	GG18746 [Drosophila erecta]	1,00E-18
AM855149	1,00E-06		
		(sp Q9H8W5) Tripartite motif-containing protein 45 OS=Homo sapiens GN=TRIM45	7,00E-12
AM855156	1,00E-06		
AM855201	1,00E-06		
		(sp Q9CSH3) Exosome complex exonuclease RRP44 OS=Mus musculus GN=Dis3 SV=4	1,00E-49
AM855209	1,00E-06		

AM855217	1,00E-05	(sp Q8BT18) Serine/arginine repetitive matrix protein 2 OS=Mus musculus GN=Srrm2	4,00E-48
AM855218	1,00E-06	(sp O08605) MAP kinase-interacting serine/threonine-protein kinase 1 OS=Mus musculus GN=Mknk1	1,00E-112
AM855237	1,00E-06	(sp Q8NWC4) Threonine dehydratase catabolic OS=Staphylococcus aureus (strain MW2) GN=tdcB	4,00E-18
AM855252	1,00E-06	(sp Q4R4R0) 26S protease regulatory subunit 7 OS=Macaca fascicularis GN=PSMC2	0,00E+00
AM855306	1,00E-06		
AM855307	1,00E-06		
AM855318	1,00E-05	PREDICTED: hypothetical protein [Gallus gallus]	3,00E-09
AM855329	1,00E-06	(sp AOJM12) Multiple epidermal growth factor-like domains 10 OS=Xenopus tropicalis GN=megf10	8,00E-06
AM855348	1,00E-06		
AM855362	1,00E-05		
AM855367	1,00E-06		
AM855375	1,00E-06		
AM855376	1,00E-05	PREDICTED: hypothetical protein [Danio rerio]	9,00E-17
AM855378	1,00E-06		
AM855384	1,00E-06		
AM855386	1,00E-06	(sp Q8WV99) AN1-type zinc finger protein 2B OS=Homo sapiens GN=ZFAND2B	5,00E-51
AM855396	1,00E-06		
AM855397	1,00E-06		
AM855402	1,00E-06	(sp Q6DFN1) NADH dehydrogenase [ubiquinone] 1 alpha subcomplex assembly factor 3 OS=Xenopus tropicalis	1,00E-24
AM855404	1,00E-06	(sp Q13201) Multimerin-1 OS=Homo sapiens GN=MMRN1	7,00E-06
AM855428	1,00E-06		
AM855441	1,00E-06		
AM855462	1,00E-05		
AM855467	1,00E-06	(sp P24051) 40S ribosomal protein S27-like protein OS=Rattus norvegicus GN=Rps27l	2,00E-40
AM855480	1,00E-06		
AM855483	1,00E-06	(sp Q6NYV8) Lysocardiolipin acyltransferase 1 OS=Danio rerio GN=lclat1	4,00E-20
AM855489	1,00E-06		

AM855501	1,00E-06	(sp Q26630) 33 kDa inner dynein arm light chain, axonemal OS=Strongylocentrotus purpuratus	1,00E-111
AM855503	1,00E-05		
AM855505	1,00E-06	PREDICTED: similar to GL23246 [Hydra magnipapillata]	2,00E-06
AM855520	1,00E-06		
AM855521	1,00E-05	(sp A8XMW6) Cell death abnormality protein 1 OS=Caenorhabditis briggsae GN=ced-1	6,00E-06
AM855522	1,00E-05	(sp P06155) Integrase OS=Enterobacteria phage phi80 GN=int	1,00E-85
AM855534	1,00E-06		
AM855621	1,00E-06		
AM855622	1,00E-06	(sp Q9NL98) Peroxiredoxin OS=Ascaris suum	3,00E-80
AM855629	1,00E-06		
AM855636	1,00E-06		
AM855638	1,00E-06	(sp O70277) Tripartite motif-containing protein 3 OS=Rattus norvegicus GN=Trim3	2,00E-06
AM855641	1,00E-05	(sp Q5VYJ5) MAM and LDL-receptor class A domain-containing protein C10orf112 OS=Homo sapiens	1,00E-36
AM855648	1,00E-05		
AM855659	1,00E-06	(sp Q5RD12) Heat shock factor-binding protein 1 OS=Pongo abelii GN=HSBP1	5,00E-20
AM855678	1,00E-06	(sp Q53T59) HCLS1-binding protein 3 OS=Homo sapiens GN=HS1BP3	3,00E-10
AM855679	1,00E-06		
AM855683	1,00E-05	(sp P27463) Retinal dehydrogenase 1 OS=Gallus gallus GN=ALDH1A1	1,00E-55
AM855689	1,00E-06		
AM855690	1,00E-06		
AM855708	1,00E-06	(sp Q03412) Innexin unc-7 OS=Caenorhabditis elegans GN=unc-7	8,00E-28
AM855731	1,00E-06		
AM855741	1,00E-06	(sp O54879) High mobility group protein B3 OS=Mus musculus GN=Hmgb3	1,00E-27
AM855744	1,00E-06		
AM855759	1,00E-06		
AM855766	1,00E-06		
AM855774	1,00E-05		
AM855777	1,00E-06	(sp Q8R1X6) Spartin OS=Mus musculus GN=Spg20	5,00E-06

AM855779	1,00E-05	(sp P58545) BTB/POZ domain-containing protein 3 OS=Mus musculus GN=Btbd3	2,00E-40
AM855785	1,00E-05	(sp Q5ZM98) Stress-70 protein, mitochondrial OS=Gallus gallus GN=HSPA9 (sp Q68FW6) Uncharacterized protein C14orf50 homolog OS=Rattus norvegicus	0,00E+00 6,00E-30
AM855799	1,00E-05		
AM855801	1,00E-06		
AM855811	1,00E-06	(sp Q6DDB9) RNA-binding protein PNO1 OS=Xenopus tropicalis GN=pno1	3,00E-96
AM855817	1,00E-06		
AM855826	1,00E-06	(sp P17710) Hexokinase-1 OS=Mus musculus GN=Hk1	1,00E-116
AM855832	1,00E-06		
AM855843	1,00E-05		
AM855848	1,00E-05	(sp Q6PDJ1) VWFA and cache domain-containing protein 1 OS=Mus musculus GN=Cachd1	4,00E-12
AM855853	1,00E-06	(sp Q6XVN8) Microtubule-associated proteins 1A/1B light chain 3A OS=Rattus norvegicus GN=Map1lc3a	2,00E-46
AM855861	1,00E-06	(sp Q8HXD5) RING finger protein 220 OS=Macaca fascicularis GN=RNF220	1,00E-13
AM855871	1,00E-06	(sp Q8BGN3) Ectonucleotide pyrophosphatase/phosphodiesterase family member 6 OS=Mus musculus GN=Enpp6	8,00E-06
AM855872	1,00E-06	(sp O75351) Vacuolar protein sorting-associated protein 4B OS=Homo sapiens GN=VPS4B	4,00E-98
AM855884	1,00E-06		
AM855888	1,00E-06	hypothetical protein BRAFLDRAFT_86827 [Branchiostoma floridae]	1,00E-46
AM855891	1,00E-06		
AM855900	1,00E-06		
AM855909	1,00E-05	(sp Q6PCB7) Long-chain fatty acid transport protein 1 OS=Homo sapiens GN=SLC27A1	8,00E-30
AM855915	1,00E-06		
AM855921	1,00E-06		
AM855924	1,00E-06		
AM855926	1,00E-06		
AM855941	1,00E-06	(sp Q9CPW4) Actin-related protein 2/3 complex subunit 5 OS=Mus musculus GN=Arpc5	4,00E-38
AM855959	1,00E-06		

AM855974	1,00E-06		
AM855984	1,00E-06		
AM856009	1,00E-06	(sp Q5X106) Probable histone acetyltransferase MYST1 OS=Rattus norvegicus GN=Myst1	2,00E-36
AM856010	1,00E-06	(sp Q9D0M3) Cytochrome c1, heme protein, mitochondrial OS=Mus musculus GN=Cyc1	6,00E-70
AM856015	1,00E-06		
AM856043	1,00E-06		
AM856044	1,00E-05	(sp Q8N7M0) Tctex1 domain-containing protein 1 OS=Homo sapiens GN=TCTEX1D1	1,00E-17
AM856048	1,00E-06		
AM856053	1,00E-06	(sp P54213) Caltractin OS=Dunaliella salina	6,00E-32
AM856067	1,00E-06	(sp Q9R0Q8) C-type lectin domain family 4 member E OS=Mus musculus GN=Clec4e	1,00E-06
AM856069	1,00E-06		
AM856079	1,00E-06	(sp Q8K2C6) NAD-dependent deacetylase sirtuin-5 OS=Mus musculus GN=Sirt5	2,00E-76
AM856080	1,00E-06		
AM856087	1,00E-06	(sp Q6DHL5) Leucine-rich repeat-containing protein 57 OS=Danio rerio GN=Irrc57	6,00E-66
AM856089	1,00E-05	(sp P12107) Collagen alpha-1(XI) chain OS=Homo sapiens GN=COL11A1 SV=4	7,00E-40
AM856116	1,00E-06		
AM856121	1,00E-06		
AM856123	1,00E-05		
AM856143	1,00E-06		
AM856145	1,00E-06		
AM856147	1,00E-06		
AM856153	1,00E-06	(sp P35479) Leukocyte cysteine proteinase inhibitor 1 OS=Sus scrofa (sp Q24157) Beta-1,3-galactosyltransferase brn OS=Drosophila melanogaster GN=brn	7,00E-07
AM856163	1,00E-06		4,00E-14
AM856169	1,00E-06		
AM856175	1,00E-06		
AM856176	1,00E-06		
AM856188	1,00E-06		

AM856189	1,00E-06	(sp Q16RX1) Mediator of RNA polymerase II transcription subunit 21 OS=Aedes aegypti GN=MED21	3,00E-43
AM856192	1,00E-06		
AM856198	1,00E-06		
AM856199	1,00E-06		
AM856208	1,00E-06		
AM856210	1,00E-06		
AM856211	1,00E-06	(sp Q95NR9) Calmodulin OS=Metridium senile (sp Q29HY3) Cdc42 homolog OS=Drosophila pseudoobscura pseudoobscura GN=Cdc42	1,00E-77
AM856213	1,00E-06		1,00E-94
AM856220	1,00E-06	(sp A1A4L6) Protein C-ets-2 OS=Bos taurus GN=ETS2	2,00E-52
AM856233	1,00E-06		
AM856257	1,00E-06		
AM856264	1,00E-06		
AM856277	1,00E-05	(sp Q9NL98) Peroxiredoxin OS=Ascaris suum	3,00E-80
AM856290	1,00E-06		
AM856305	1,00E-05		
AM856306	1,00E-06	(sp P61917) Epididymal secretory protein E1 OS=Pan troglodytes GN=NPC2 (sp Q7SXB3) Leucine-rich repeat-containing protein 40 OS=Danio rerio GN=Irrc40	3,00E-24
AM856313	1,00E-06		7,00E-09
AM856320	1,00E-06	(sp Q9X6U2) D-beta-hydroxybutyrate dehydrogenase OS=Ralstonia eutropha	3,00E-27
AM856323	1,00E-06		
AM856330	1,00E-06		
AM856335	1,00E-06		
AM856336	1,00E-06		
AM856343	1,00E-06		
AM856348	1,00E-06	(sp Q17QE5) Calcium and integrin-binding protein 1 OS=Bos taurus GN=CIB1	4,00E-37
AM856355	1,00E-06		
AM856356	1,00E-06		
AM856360	1,00E-06		
AM856365	1,00E-06		
AM856372	1,00E-05		

AM856374	1,00E-06	(sp Q7T3H9) Putative hydroxypyruvate isomerase OS=Danio rerio GN=hyi	3,00E-70
AM856380	1,00E-05		
AM856384	1,00E-06		
AM856393	1,00E-06		
AM856394	1,00E-06		
AM856398	1,00E-06	(sp Q5R4U9) Sorcin OS=Pongo abelii GN=SRI	5,00E-24
AM856410	1,00E-05		
AM856417	1,00E-06		
AM856419	1,00E-06	(sp P55780) NADH-ubiquinone oxidoreductase chain 2 OS=Gadus morhua GN=MT-ND2	5,00E-11
AM856435	1,00E-05	(sp Q5T011) Uncharacterized protein KIAA0467 OS=Homo sapiens GN=KIAA0467	9,00E-09
AM856440	1,00E-06		
AM856446	1,00E-06		
AM856458	1,00E-06	predicted protein [Nematostella vectensis]	2,00E-11
AM856472	1,00E-05		
AM856474	1,00E-05	(sp P18503) Short-chain collagen C4 (Fragment) OS=Ephydatia muelleri	3,00E-09
AM856491	1,00E-06		
AM856520	1,00E-05	(sp Q9D3E6) Cohesin subunit SA-1 OS=Mus musculus GN=Stag1	3,00E-42
AM856554	1,00E-06	hypothetical protein BRAFLDRAFT_100697 [Branchiostoma floridae]	8,00E-08
AM856561	1,00E-05	(sp Q5ZMP6) AP-2 complex subunit mu-1 OS=Gallus gallus GN=AP2M1	0,00E+00
AM856564	1,00E-06		
AM856566	1,00E-05		
AM856577	1,00E-06	(sp Q6AZB8) Putative nuclease HARBI1 OS=Danio rerio GN=harbi1	5,00E-29
AM856582	1,00E-05		
AM856590	1,00E-06	(sp P54145) Putative ammonium transporter 1 OS=Caenorhabditis elegans GN=amt-1	3,00E-27
AM856597	1,00E-05	(sp Q9Y6W6) Dual specificity protein phosphatase 10 OS=Homo sapiens GN=DUSP10	3,00E-86
AM856598	1,00E-06	(sp A0JPB5) Membrane protein FAM174 OS=Xenopus tropicalis GN=fam174	5,00E-08
AM856611	1,00E-06		
AM856619	1,00E-06		
AM856620	1,00E-06	(sp Q6TAW2) Stress-associated endoplasmic reticulum protein 2 OS=Mus musculus GN=Serp2	3,00E-23

AM856621	1,00E-06	(sp P34611) B-box type zinc finger protein ncl-1 OS=Caenorhabditis elegans GN=ncl-1	3,00E-07
AM856631	1,00E-06		
AM856646	1,00E-05		
AM856653	1,00E-05	(sp Q9VA11) Probable complex I intermediate-associated protein 30, mitochondrial OS=Drosophila melanogaster	5,00E-39
AM856658	1,00E-06	(sp Q4S3C1) UPF0402 protein OS=Tetraodon nigroviridis GN=GSTENG00024711001	8,00E-24
AM856664	1,00E-06		
AM856665	1,00E-06		
AM856671	1,00E-06	(sp Q96MM6) Heat shock 70 kDa protein 12B OS=Homo sapiens GN=HSPA12B	1,00E-49
AM856674	1,00E-06		
AM856678	1,00E-06	(sp Q28735) Transmembrane emp24 domain-containing protein 10 OS=Oryctolagus cuniculus	8,00E-24
AM856684	1,00E-06		
AM856695	1,00E-06	(sp Q9UKA2) F-box/LRR-repeat protein 4 OS=Homo sapiens GN=FBXL4	1,00E-06
AM856701	1,00E-06		
AM856704	1,00E-06	(sp Q63625) PHD and RING finger domain-containing protein 1 OS=Rattus norvegicus GN=Phrf1	1,00E-23
AM856707	1,00E-06		
AM856711	1,00E-06	(sp Q96MM6) Heat shock 70 kDa protein 12B OS=Homo sapiens GN=HSPA12B	2,00E-40
AM856722	1,00E-06	(sp P53569) CCAAT/enhancer-binding protein zeta OS=Mus musculus GN=Cebpz	4,00E-09
AM856723	1,00E-06	(sp Q28G94) Dynein light chain 1, axonemal OS=Xenopus tropicalis GN=dnal1	1,00E-76
AM856724	1,00E-06		
AM856725	1,00E-05		
AM856729	1,00E-05	(sp P62840) Ubiquitin-conjugating enzyme E2 D2 OS=Xenopus laevis GN=ube2d2	8,00E-42
AM856730	1,00E-05	(sp Q3SXD3) HD domain-containing protein 2 OS=Mus musculus GN=Hddc2	5,00E-59
AM856750	1,00E-06		
AM856754	1,00E-06		

AM856759	1,00E-06		
AM856777	1,00E-06	(sp Q86VS8) Protein Hook homolog 3 OS=Homo sapiens GN=HOOK3	5,00E-64
AM856783	1,00E-06		
AM856807	1,00E-06	(sp P04352) Calmodulin OS=Chlamydomonas reinhardtii	4,00E-46
		(sp Q96B70) Leukocyte receptor cluster member 9 OS=Homo sapiens	
AM856814	1,00E-06	GN=LENG9	2,00E-17
AM856819	1,00E-06		
AM856825	1,00E-06		
		(sp Q66H03) Uncharacterized protein LOC285141 homolog OS=Rattus norvegicus	
AM856826	1,00E-05		7,00E-06
AM856833	1,00E-06		
AM856841	1,00E-06	(sp A3KNS9) Apoptosis-related protein 3 OS=Danio rerio GN=apr3	7,00E-24
AM856858	1,00E-06		
AM856868	1,00E-06		
AM856885	1,00E-06		
		(sp Q0VFN8) Uncharacterized protein C9orf117 homolog OS=Xenopus tropicalis	
AM856886	1,00E-06		1,00E-77
AM856895	1,00E-06		
AM856899	1,00E-06		
AM856919	1,00E-06		
		(sp Q91X20) Set1/Ash2 histone methyltransferase complex subunit ASH2 OS=Mus musculus	
AM856921	1,00E-06		3,00E-48
AM856923	1,00E-05		
AM856932	1,00E-06		
AM856933	1,00E-06		
AM856937	1,00E-06		
AM856945	1,00E-06		
AM856948	1,00E-06	(sp Q16956) 78 kDa glucose-regulated protein OS=Aplysia californica	2,00E-24
AM856951	1,00E-05		
AM856952	1,00E-05	(sp Q8CFC2) Myelin transcription factor 1 OS=Mus musculus GN=Myt1	9,00E-07
AM856956	1,00E-06	(sp A5PK63) 40S ribosomal protein S17 OS=Bos taurus GN=RPS17	1,00E-55
AM856957	1,00E-06	predicted protein [Nematostella vectensis]	3,00E-06
AM856959	1,00E-06	(sp Q96F15) GTPase IMAP family member 5 OS=Homo sapiens GN=GIMAP5	2,00E-08
AM856969	1,00E-06	(sp P58743) Prestin OS=Homo sapiens GN=SLC26A5	5,00E-32

AM856972	1,00E-06	(sp P50580) Proliferation-associated protein 2G4 OS=Mus musculus GN=Pa2g4	1,00E-100
AM856979	1,00E-06	(sp Q9CZ30) Obg-like ATPase 1 OS=Mus musculus GN=Ola1	1,00E-110
AM856984	1,00E-06	(sp Q969Q1) E3 ubiquitin-protein ligase TRIM63 OS=Homo sapiens GN=TRIM63	4,00E-06
AM856990	1,00E-06	(sp Q0VE82) Copine-7 OS=Mus musculus GN=Cpne7	1,00E-13
AM857016	1,00E-05	(sp Q9Y6G5) COMM domain-containing protein 10 OS=Homo sapiens GN=COMMD10	5,00E-39
AM857034	1,00E-06		
AM857046	1,00E-06		
AM857047	1,00E-06		
AM857049	1,00E-06		
AM857059	1,00E-06	(sp Q502K9) Selenoprotein T2 OS=Danio rerio GN=selt2	3,00E-45
AM857068	1,00E-06		
AM857075	1,00E-06	(sp Q17RR3) Pancreatic lipase-related protein 3 OS=Homo sapiens GN=PNLIPRP3	7,00E-09
AM857081	1,00E-06	GH23959 [Drosophila grimshawi]	1,00E-08
AM857097	1,00E-06	(sp Q9CZJ2) Heat shock 70 kDa protein 12B OS=Mus musculus GN=Hspa12b	2,00E-36
AM857099	1,00E-06		
AM857106	1,00E-05		
AM857110	1,00E-05	(sp B8JMH0) Putative transferase C1orf69 homolog, mitochondrial OS=Danio rerio GN=ch1073-162b20.1	7,00E-18
AM857112	1,00E-06		
AM857116	1,00E-06	(sp Q61D18) Laccase-10 OS=Arabidopsis thaliana GN=LAC10	2,00E-16
AM857121	1,00E-06	(sp Q5ZJL9) SAM domain and HD domain-containing protein 1 OS=Gallus gallus GN=SAMHD1	6,00E-42
AM857140	1,00E-06		
AM857141	1,00E-06	(sp Q2YGT9) 60S ribosomal protein L6 OS=Sus scrofa GN=RPL6	2,00E-64
AM857154	1,00E-06		
AM857157	1,00E-06		
AM857160	1,00E-06	(sp Q8BLJ3) PI-PLC X domain-containing protein 3 OS=Mus musculus GN=Plcxd3	4,00E-35
AM857161	1,00E-05	(sp Q2KHW8) Gem-associated protein 6 OS=Bos taurus GN=GEMIN6	2,00E-22
AM857162	1,00E-06		

AM857172	1,00E-06		
AM857173	1,00E-06	(sp Q9Z2K0) Death effector domain-containing protein OS=Rattus norvegicus GN=Dedd	9,00E-25
AM857177	1,00E-05		
AM857182	1,00E-06	(sp Q761C5) Pyroglutamyl-peptidase 1 OS=Rattus norvegicus GN=Pgpep1	7,00E-55
AM857183	1,00E-06		
AM857184	1,00E-06	(sp P61354) 60S ribosomal protein L27 OS=Rattus norvegicus GN=Rpl27	5,00E-53
AM857195	1,00E-06		
AM857199	1,00E-06		
AM857200	1,00E-06		
AM857210	1,00E-05		
AM857211	1,00E-06	(sp Q5ZMM5) Zinc finger protein 706 OS=Gallus gallus GN=ZNF706 PE=4 (sp Q9R016) Baculoviral IAP repeat-containing protein 4 OS=Rattus norvegicus GN=Xiap	1,00E-25 6,00E-06
AM857221	1,00E-06		
AM857225	1,00E-06	(sp Q64430) Copper-transporting ATPase 1 OS=Mus musculus GN=Atp7a	1,00E-64
AM857237	1,00E-06		
AM857240	1,00E-06		
AM857249	1,00E-06	(sp Q6TFL4) Kelch-like protein 24 OS=Homo sapiens GN=KLHL24 (sp P32198) Carnitine O-palmitoyltransferase 1, liver isoform OS=Rattus norvegicus GN=Cpt1a	2,00E-24 2,00E-17
AM857294	1,00E-06		
AM857297	1,00E-06		
AM857299	1,00E-06	(sp Q801P0) LIM domain transcription factor LMO4-B OS=Xenopus laevis GN=Imo4-B (sp Q21086) Guanine nucleotide-binding protein-like 3 homolog OS=Caenorhabditis elegans GN=nst-1	2,00E-44 1,00E-45
AM857302	1,00E-06		
AM857308	1,00E-06		
AM857313	1,00E-05		
AM857315	1,00E-06		
AM857327	1,00E-06	(sp Q969S6) Transmembrane protein 203 OS=Homo sapiens GN=TMEM203	2,00E-11
AM857335	1,00E-06	(sp Q96RW7) Hemicentin-1 OS=Homo sapiens GN=HMCN1	2,00E-29
AM857356	1,00E-06		
AM857357	1,00E-06	(sp P21895) Histone H1 OS=Chironomus thummi thummi	7,00E-12
AM857375	1,00E-05		

AM857381	1,00E-05	(sp P49724) Ornithine aminotransferase, mitochondrial OS=Drosophila ananassae GN=Oat	3,00E-29
AM857382	1,00E-06		
AM857387	1,00E-06	(sp Q61941) NAD(P) transhydrogenase, mitochondrial OS=Mus musculus GN=Nnt	2,00E-54
AM857389	1,00E-06		
AM857392	1,00E-06	(sp Q5R889) Putative adenosylhomocysteinase 3 OS=Pongo abelii GN=AHCYL2	6,00E-46
AM857401	1,00E-06	(sp Q15024) Exosome complex exonuclease RRP42 OS=Homo sapiens GN=EXOSC7	6,00E-89
AM857405	1,00E-06	(sp Q9D752) Mitotic spindle assembly checkpoint protein MAD2B OS=Mus musculus GN=Mad2l2	2,00E-43
AM857412	1,00E-05	(sp P07290) Myosin, essential light chain, adductor muscle OS=Patinopecten yessoensis	2,00E-45
AM857415	1,00E-06		
AM857417	1,00E-05	(sp Q9XT62) Cell cycle checkpoint protein RAD17 OS=Cercopithecus aethiops GN=RAD17	1,00E-17
AM857426	1,00E-06		
AM857429	1,00E-06	(sp Q6EE31) T-complex protein 1 subunit theta OS=Gallus gallus GN=CCT8	0,00E+00
AM857435	1,00E-06		
AM857436	1,00E-06		
AM857440	1,00E-06		
AM857447	1,00E-06		
AM857451	1,00E-06		
AM857461	1,00E-06	(sp A6H7E2) Cyclin-dependent kinase 2-interacting protein OS=Bos taurus GN=CINP	7,00E-14
AM857472	1,00E-06		
AM857473	1,00E-06	(sp Q9BDK2) Allograft inflammatory factor 1 OS=Bos taurus GN=AIF1	1,00E-33
AM857474	1,00E-06		
AM857486	1,00E-06	(sp Q58CY6) Uncharacterized protein C1orf93 homolog OS=Bos taurus	6,00E-41
AM857494	1,00E-05		
AM857505	1,00E-06		
AM857506	1,00E-06		
AM857541	1,00E-06		

AM857553	1,00E-06		
AM857557	1,00E-06	(sp P24051) 40S ribosomal protein S27-like protein OS=Rattus norvegicus	
AM857563	1,00E-06	GN=Rps271	2,00E-40
AM857580	1,00E-06	(sp Q0V FH6) 39S ribosomal protein L22, mitochondrial OS=Xenopus tropicalis	
AM857592	1,00E-06	GN=mrpl22	1,00E-31
AM857594	1,00E-06	(sp Q86Y82) Syntaxin-12 OS=Homo sapiens GN=STX12	2,00E-24
AM857605	1,00E-06	(sp Q5M7G4) Putative adenylate kinase-like protein C9orf98 homolog OS=Xenopus laevis	4,00E-81
AM857622	1,00E-06	(sp Q5RD88) Lysine-specific demethylase 4A OS=Pongo abelii GN=KDM4A	7,00E-12
AM857623	1,00E-05		
AM857625	1,00E-05		
AM857627	1,00E-06	(sp Q2KIU7) Radial spoke head protein 9 homolog OS=Bos taurus GN=RSPH9	5,00E-81
AM857635	1,00E-05	(sp P62317) Small nuclear ribonucleoprotein Sm D2 OS=Mus musculus GN=Snrpd2	2,00E-44
AM857640	1,00E-06	(sp Q28DG7) Ubiquitin-associated domain-containing protein 1 OS=Xenopus tropicalis GN=ubac1	3,00E-35
AM857641	1,00E-06	(sp A2VE44) Mediator of RNA polymerase II transcription subunit 25 OS=Bos taurus GN=MED25	6,00E-33
AM857658	1,00E-06		
AM857659	1,00E-06		
AM857661	1,00E-06	(sp Q5ZLC7) Microtubule-associated protein RP/EB family member 1 OS=Gallus gallus GN=MAPRE1	2,00E-55
AM857662	1,00E-06		
AM857682	1,00E-06		
AM857687	1,00E-06		
AM857692	1,00E-06		
AM857695	1,00E-06	(sp Q58CY6) Uncharacterized protein C1orf93 homolog OS=Bos taurus	6,00E-41
AM857696	1,00E-06		
AM857700	1,00E-06		
AM857703	1,00E-06		

AM857714	1,00E-06	(sp Q55FM5) Myotrophin homolog OS=Dictyostelium discoideum GN=mtpn	2,00E-27
AM857721	1,00E-06		
AM857725	1,00E-06		
AM857726	1,00E-06		
AM857737	1,00E-06	(sp Q9BT43) DNA-directed RNA polymerase III subunit RPC7-like OS=Homo sapiens GN=POLR3GL	2,00E-32
AM857738	1,00E-05		
AM857751	1,00E-06		
AM857760	1,00E-06	(sp P08228) Superoxide dismutase [Cu-Zn] OS=Mus musculus GN=Sod1	1,00E-59
AM857766	1,00E-05	(sp P51958) Cell division control protein 2 homolog OS=Carassius auratus GN=cdc2	1,00E-139
AM857767	1,00E-06		
AM857768	1,00E-06	(sp P26270) 26S proteasome non-ATPase regulatory subunit 7 OS=Drosophila melanogaster GN=Mov34 SV=6	1,00E-118
AM857769	1,00E-06	(sp P42765) 3-ketoacyl-CoA thiolase, mitochondrial OS=Homo sapiens GN=ACAA2	6,00E-87
AM857771	1,00E-06		
AM857782	1,00E-06		
AM857783	1,00E-06	(sp Q9VJ38) 39S ribosomal protein L13, mitochondrial OS=Drosophila melanogaster GN=mRpL13	2,00E-43
AM857787	1,00E-06	(sp Q6DER2) 60S ribosomal protein L36 OS=Xenopus tropicalis GN=rpl36	2,00E-31
AM857796	1,00E-05		
AM857806	1,00E-06	(sp A9UMG5) Protein IMPACT-B OS=Xenopus tropicalis GN=impact-B	1,00E-63
AM857811	1,00E-06	(sp Q6NRK9) Cyclin-related protein FAM58A OS=Xenopus laevis GN=fam58a	1,00E-60
AM857817	1,00E-06		
AM857818	1,00E-06	(sp Q9CR61) NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 7 OS=Mus musculus	4,00E-18
AM857827	1,00E-06		
AM857829	1,00E-06		
AM857835	1,00E-06		
AM857843	1,00E-06		
AM857844	1,00E-06		
AM857845	1,00E-06	predicted protein [Nematostella vectensis]	3,00E-06

AM857851	1,00E-06	(sp O69060) Probable alpha-ketoglutarate-dependent hypophosphite dioxxygenase OS=Pseudomonas stutzeri	1,00E-09
AM857861	1,00E-06		
AM857866	1,00E-06	(sp Q5ZIP6) OTU domain-containing protein 6B OS=Gallus gallus	4,00E-06
AM857897	1,00E-06	GN=OTUD6B	
AM857898	1,00E-06		
AM857899	1,00E-06	(sp Q9CQY6) Uncharacterized protein C6orf125 homolog OS=Mus musculus	5,00E-10
AM857911	1,00E-06		
AM857916	1,00E-06	(sp Q9ESN6) Tripartite motif-containing protein 2 OS=Mus musculus	4,00E-06
AM857924	1,00E-05	GN=Trim2	2,00E-60
AM857928	1,00E-06	(sp P17712) Glucokinase OS=Rattus norvegicus GN=Gck	8,00E-43
AM857933	1,00E-06		
AM857937	1,00E-05	(sp Q4G0N4) UPF0465 protein C5orf33 OS=Homo sapiens GN=C5orf33	2,00E-26
AM857938	1,00E-06		
AM857939	1,00E-06		
AM857941	1,00E-06		
AM857945	1,00E-06	(sp Q05AK9) Coiled-coil domain-containing protein 72 OS=Danio rerio	3,00E-10
AM857948	1,00E-06	GN=ccdc72	
AM857963	1,00E-06		
AM857973	1,00E-06		
AM857988	1,00E-06		
AM857996	1,00E-06	(sp Q9Y547) Heat shock protein beta-11 OS=Homo sapiens GN=HSPB11	6,00E-31
AM858009	1,00E-06	(sp P62876) DNA-directed RNA polymerases I, II, and III subunit RPABC5	2,00E-32
AM858018	1,00E-06	OS=Mus musculus GN=Polr2l	2,00E-47
AM858019	1,00E-06	(sp Q5U3J8) Protein LTV1 homolog OS=Danio rerio GN=ltv1	3,00E-78
AM858020	1,00E-05		
AM858050	1,00E-06	(sp Q7T3E4) Lariat debranching enzyme OS=Danio rerio GN=dbr1	2,00E-15
		(sp Q5R5L7) Arrestin domain-containing protein 3 OS=Pongo abelii	
		GN=ARRDC3	

AM858060	1,00E-06	(sp Q9Z219) Succinyl-CoA ligase [ADP-forming] subunit beta, mitochondrial OS=Mus musculus GN=Sucla2	2,00E-86
AM858070	1,00E-05	(sp O88794) Pyridoxine-5'-phosphate oxidase OS=Rattus norvegicus GN=Pnpo	5,00E-53
AM858094	1,00E-06	(sp Q7YR30) Protein phosphatase 1 regulatory subunit 11 OS=Pan troglodytes GN=PPP1R11	5,00E-14
AM858119	1,00E-06	(sp A6QLI0) Mammalian ependymin-related protein 1 OS=Bos taurus GN=EPDR1	9,00E-06
AM858132	1,00E-05	(sp Q5RE11) Tetraspanin-3 OS=Pongo abelii GN=TSPAN3	1,00E-06
AM858144	1,00E-06	(sp Q6DIB5) Multiple epidermal growth factor-like domains 10 OS=Mus musculus GN=Megf10	1,00E-06
AM858155	1,00E-06	(sp Q1HG43) Dual oxidase maturation factor 1 OS=Homo sapiens GN=DUOXA1	6,00E-20
AM858156	1,00E-06	(sp O43300) Leucine-rich repeat transmembrane neuronal protein 2 OS=Homo sapiens GN=LRRTM2	1,00E-11
AM858175	1,00E-05	(sp Q5BJP5) UPF0414 transmembrane protein C20orf30 homolog OS=Rattus norvegicus	5,00E-35
AM858179	1,00E-06		
AM858185	1,00E-06		
AM858189	1,00E-05	(sp Q9NYG2) Palmitoyltransferase ZDHHC3 OS=Homo sapiens GN=ZDHHC3	8,00E-23
AM858191	1,00E-06	(sp Q96BK5) Pin2-interacting protein X1 OS=Homo sapiens GN=PINX1	9,00E-34
AM858193	1,00E-06	(sp Q24307) Apoptosis 2 inhibitor OS=Drosophila melanogaster GN=lap2	3,00E-17
AM858212	1,00E-06		
AM858215	1,00E-06	(sp Q3T0B6) Complement component 1 Q subcomponent-binding protein, mitochondrial OS=Bos taurus GN=C1QBP	1,00E-15
AM858216	1,00E-06	(sp P49395) 40S ribosomal protein S3a OS=Aplysia californica GN=RPS3A	1,00E-107
AM858218	1,00E-06	(sp Q96MD2) UPF0536 protein C12orf66 OS=Homo sapiens GN=C12orf66	2,00E-82
AM858226	1,00E-06	(sp Q9UN37) Vacuolar protein sorting-associated protein 4A OS=Homo sapiens GN=VPS4A	2,00E-91
AM858229	1,00E-05	(sp Q4R642) T-complex-associated testis-expressed protein 1 OS=Macaca fascicularis GN=TCTE1	1,00E-50
AM858233	1,00E-06		
AM858235	1,00E-06		

AM858239	1,00E-05		
AM858244	1,00E-06	(sp Q7SYJ9) Sperm-associated antigen 7 homolog OS=Danio rerio GN=spag7	1,00E-50
AM858246	1,00E-05	(sp Q5W5X9) Tetratricopeptide repeat protein 23 OS=Homo sapiens	4,00E-26
AM858247	1,00E-06	GN=TTC23	
AM858248	1,00E-06		
AM858251	1,00E-06	(sp P60517) Gamma-aminobutyric acid receptor-associated protein	
AM858255	1,00E-05	OS=Rattus norvegicus GN=Gabarap	1,00E-58
AM858257	1,00E-06	(sp Q6NRW3) Kinetochores protein Spc24 OS=Xenopus laevis GN=spc24	3,00E-09
AM858259	1,00E-06	(sp P62925) Eukaryotic translation initiation factor 5A OS=Spodoptera	
AM858289	1,00E-05	frugiperda GN=eIF-5A	9,00E-52
AM858295	1,00E-06	(sp Q15013) MAD2L1-binding protein OS=Homo sapiens GN=MAD2L1BP	6,00E-15
AM858296	1,00E-06	(sp Q9D0M3) Cytochrome c1, heme protein, mitochondrial OS=Mus musculus	
AM858303	1,00E-06	GN=Cyc1	6,00E-70
AM858311	1,00E-06	(sp Q3SZJ4) Prostaglandin reductase 1 OS=Bos taurus GN=PTGR1	7,00E-57
AM858312	1,00E-05		
AM858325	1,00E-06		
AM858331	1,00E-06		
AM858332	1,00E-06	(sp P09954) Carboxypeptidase A1 OS=Sus scrofa GN=CPA1	1,00E-33
AM858342	1,00E-05	(sp P30044) Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5	5,00E-52
AM858350	1,00E-06	(sp P61801) Aprataxin OS=Xenopus tropicalis GN=aptx	3,00E-15
AM858352	1,00E-06	(sp P06623) 2',3'-cyclic-nucleotide 3'-phosphodiesterase OS=Bos taurus	
AM858382	1,00E-06	GN=CNP	7,00E-23
AM858384	1,00E-06	(sp Q03591) Complement factor H-related protein 1 OS=Homo sapiens	
AM858394	1,00E-06	GN=CFHR1	5,00E-06
AM858413	1,00E-06	glucosidase [Chryseobacterium gleum ATCC 35910]	4,00E-14
AM858426	1,00E-06	(sp Q05062) Cell division control protein 42 homolog OS=Caenorhabditis	
		elegans GN=cdc-42	2,00E-38

AM858429	1,00E-06		
AM858445	1,00E-06		
AM858446	1,00E-05	(sp Q9VVN2) Probable 28S ribosomal protein S26, mitochondrial OS=Drosophila melanogaster GN=mRpS26	4,00E-13
AM858450	1,00E-06		
AM858456	1,00E-05		
AM858458	1,00E-06		
AM858461	1,00E-06	PREDICTED: similar to ENSANGP00000017748, partial [Strongylocentrotus purpuratus]	2,00E-15
AM858467	1,00E-06	(sp Q9CR80) Protein FAM32A OS=Mus musculus GN=Fam32a	6,00E-09
AM858514	1,00E-06		
AM858537	1,00E-06	(sp P55068) Brevican core protein OS=Rattus norvegicus GN=Bcan	9,00E-07
AM858547	1,00E-05	(sp Q95ND5) Caspase-3 OS=Sus scrofa GN=CASP3	4,00E-06
AM858556	1,00E-06		
AM858569	1,00E-05	(sp P55210) Caspase-7 OS=Homo sapiens GN=CASP7	2,00E-16
AM858581	1,00E-06	(sp P40692) DNA mismatch repair protein Mlh1 OS=Homo sapiens GN=MLH1 (sp Q8VE65) Transcription initiation factor TFIID subunit 12 OS=Mus musculus GN=Taf12	4,00E-84
AM858583	1,00E-05		2,00E-44
AM858606	1,00E-05		
AM858607	1,00E-06		
AM858608	1,00E-06	(sp Q5R613) Sorting nexin-6 OS=Pongo abelii GN=SNX6	1,00E-157
AM858609	1,00E-05		
AM858614	1,00E-05		
AM858647	1,00E-05		
AM858653	1,00E-06	(sp Q6Q311) 40S ribosomal protein S25 OS=Ovis aries GN=RPS25 (sp Q9ES30) Complement C1q tumor necrosis factor-related protein 3 OS=Mus musculus GN=C1qtnf3	4,00E-26
AM858664	1,00E-06		1,00E-06
AM858680	1,00E-06	(sp Q4R520) Endoplasmin OS=Macaca fascicularis GN=HSP90B1 (sp Q63528) Replication protein A 32 kDa subunit OS=Rattus norvegicus GN=Rpa2	0,00E+00
AM858697	1,00E-06		9,00E-07
AM858699	1,00E-06	(sp Q8TBK6) Zinc finger CCHC domain-containing protein 10 OS=Homo sapiens GN=ZCCHC10	3,00E-09
AM858701	1,00E-06	PREDICTED: hypothetical protein [Strongylocentrotus purpuratus]	1,00E-06

AM858702	1,00E-06	(sp Q2Q1W2) Tripartite motif-containing protein 71 OS=Homo sapiens GN=TRIM71	5,00E-06
AM858703	1,00E-06	(sp Q6GL69) MKI67 FHA domain-interacting nucleolar phosphoprotein-like OS=Xenopus tropicalis GN=mki67ipl	3,00E-29
AM858706	1,00E-06	(sp O18373) Selenide, water dikinase OS=Drosophila melanogaster GN=SeID	1,00E-127
AM858722	1,00E-06	PREDICTED: hypothetical protein [Danio rerio]	5,00E-13
AM858727	1,00E-06	(sp Q9UFE4) Coiled-coil domain-containing protein 39 OS=Homo sapiens GN=CCDC39	9,00E-07
AM858736	1,00E-06	(sp Q5JVL4) EF-hand domain-containing protein 1 OS=Homo sapiens GN=EFHC1	4,00E-07
AM858741	1,00E-06		
AM858743	1,00E-06	(sp P52432) DNA-directed RNA polymerases I and III subunit RPAC1 OS=Mus musculus GN=Polr1e	1,00E-92
AM858745	1,00E-05		
AM858761	1,00E-06	(sp Q9Y5J7) Mitochondrial import inner membrane translocase subunit Tim9 OS=Homo sapiens GN=TIMM9	4,00E-20
AM858762	1,00E-06		
AM858767	1,00E-06	(sp Q7SY73) Monoacylglycerol lipase abhd6-B OS=Xenopus laevis GN=abhd6- B	1,00E-48
AM858772	1,00E-06	hypothetical protein [Schistosoma mansoni]	3,00E-30
AM858773	1,00E-06	(sp Q95KV7) NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 13 OS=Bos taurus GN=NDUFA13	7,00E-19
AM858796	1,00E-06		
AM858805	1,00E-06	hypothetical protein CHLREDRAFT_115645 [Chlamydomonas reinhardtii]	2,00E-52
AM858822	1,00E-06	(sp Q9C040) Tripartite motif-containing protein 2 OS=Homo sapiens GN=TRIM2	1,00E-118
AM858828	1,00E-06		
AM858857	1,00E-06		
AM858876	1,00E-06		
AM858891	1,00E-06	(sp P46405) 40S ribosomal protein S12 OS=Sus scrofa GN=RPS12	5,00E-57
AM858895	1,00E-06		
AM858898	1,00E-06		
AM858913	1,00E-06	(sp Q5XGY9) Meiotic nuclear division protein 1 homolog OS=Xenopus laevis GN=mnd1	3,00E-57

AM858937	1,00E-06	(sp Q7RTV5) UPF0308 protein C9orf21 OS=Homo sapiens GN=C9orf21 (sp Q5F4A9) ADP-ribosylation factor-like protein 6-interacting protein 4 OS=Gallus gallus GN=ARL6IP4	1,00E-43 6,00E-18
AM858939	1,00E-06		
AM858947	1,00E-06		
AM858951	1,00E-06	PREDICTED: similar to predicted protein [Hydra magnipapillata]	5,00E-07
AM858952	1,00E-06	(sp Q8N4B1) Protein FAM109A OS=Homo sapiens GN=FAM109A	2,00E-33
AM858969	1,00E-06		
AM858974	1,00E-06		
AM858981	1,00E-06		
AM859016	1,00E-05		
AM859017	1,00E-05		
AM859025	1,00E-06		
AM859028	1,00E-06	(sp O95453) Poly(A)-specific ribonuclease PARN OS=Homo sapiens GN=PARN	2,00E-65
AM859057	1,00E-06	(sp Q9VGI8) Bloom syndrome protein homolog OS=Drosophila melanogaster GN=mus309	1,00E-15
AM859063	1,00E-05	(sp P08640) Flocculation protein FLO11 OS=Saccharomyces cerevisiae GN=MUC1	3,00E-07
AM859071	1,00E-05	(sp Q90YU6) 60S ribosomal protein L22 OS=Ictalurus punctatus GN=rpl22	3,00E-30
AM859077	1,00E-06	(sp Q96S19) UPF0585 protein C16orf13 OS=Homo sapiens GN=C16orf13	2,00E-47
AM859082	1,00E-05		
AM859089	1,00E-06	(sp P50539) MAX-interacting protein 1 OS=Homo sapiens GN=MXI1 (sp P56597) Nucleoside diphosphate kinase homolog 5 OS=Homo sapiens GN=NME5	3,00E-27 2,00E-82
AM859094	1,00E-06	(sp P20397) Nucleolin OS=Xenopus laevis GN=ncl	9,00E-07
AM859097	1,00E-06	RAD51-associated protein 1 [Salmo salar]	2,00E-07
AM859098	1,00E-06		
AM859148	1,00E-06		
AM859152	1,00E-06	hypothetical protein [Leishmania infantum] (sp P54357) Myosin-2 essential light chain OS=Drosophila melanogaster GN=Mlc-c	7,00E-06 1,00E-51
AM859167	1,00E-06	(sp Q1PRL4) Tripartite motif-containing protein 71 OS=Gallus gallus GN=TRIM71	6,00E-06
AM859173	1,00E-06	(sp P02434) 60S ribosomal protein L35a OS=Xenopus laevis GN=rpl35a	9,00E-41
AM859174	1,00E-06		

AM859184	1,00E-06	(sp Q3KRE6) Kelch domain-containing protein 2 OS=Rattus norvegicus	
AM859186	1,00E-05	GN=Klhdc2	4,00E-36
AM859189	1,00E-06	(sp O09053) Werner syndrome ATP-dependent helicase homolog OS=Mus	
AM859193	1,00E-06	musculus GN=Wrn	7,00E-10
AM859208	1,00E-06		
AM859226	1,00E-06		
AM859236	1,00E-06		
AM859241	1,00E-06		
AM859243	1,00E-06	(sp P62914) 60S ribosomal protein L11 OS=Rattus norvegicus GN=Rpl11	5,00E-83
AM859245	1,00E-05		
AM859246	1,00E-06		
AM859248	1,00E-06		
AM859250	1,00E-05		
AM859253	1,00E-06		
AM859263	1,00E-06		
AM859267	1,00E-06	(sp Q61MI6) Sulfotransferase 1C3 OS=Homo sapiens GN=SULT1C3	1,00E-25
AM859303	1,00E-06	(sp P11240) Cytochrome c oxidase subunit 5A, mitochondrial OS=Rattus	
AM859315	1,00E-06	norvegicus GN=Cox5a	3,00E-18
AM859321	1,00E-06		
AM859322	1,00E-06	(sp P51611) Host cell factor OS=Mesocricetus auratus GN=HCFC1	3,00E-10
AM859324	1,00E-06		
AM859339	1,00E-06	(sp Q5R9Z1) Vacuolar protein sorting-associated protein 29 OS=Pongo abelii	
AM859344	1,00E-05	GN=VPS29	5,00E-93
AM859353	1,00E-06		
AM859381	1,00E-05	(sp Q9FJL0) Structural maintenance of chromosomes protein 4	
AM859385	1,00E-06	OS=Arabidopsis thaliana GN=SMC4	1,00E-67
AM859386	1,00E-06	(sp P46023) G-protein coupled receptor GRL101 OS=Lymnaea stagnalis	5,00E-20
		(sp Q64373) Bcl-2-like protein 1 OS=Mus musculus GN=Bcl2l1	2,00E-14
AY496064	1,00E-06	(sp Q9VFC8) Putative glycogen [starch] synthase OS=Drosophila	
BQ426239	1,00E-06	melanogaster GN=CG6904	0,00E+00

BQ426254	1,00E-06		
BQ426259	1,00E-06	(sp Q9JLF6) Protein-glutamine gamma-glutamyltransferase K OS=Mus musculus GN=Tgm1	1,00E-39
BQ426264	1,00E-06	GA15383 [<i>Drosophila pseudoobscura pseudoobscura</i>]	2,00E-52
BQ426265	1,00E-06	(sp Q5ZMV8) Rap1 GTPase-activating protein 2 OS=Gallus gallus GN=RAP1GAP2	4,00E-06
BQ426267	1,00E-06	(sp P17710) Hexokinase-1 OS=Mus musculus GN=Hk1	1,00E-116
BQ426278	1,00E-06	(sp Q63073) Protein BTG1 OS=Rattus norvegicus GN=Btg1	5,00E-34
BQ426282	1,00E-06	(sp Q66S17) Natterin-3 OS=Thalassophryne nattereri	4,00E-06
BQ426298	1,00E-06		
BQ426316	1,00E-06		
BQ426347	1,00E-05	(sp Q9EPE9) Probable cation-transporting ATPase 13A1 OS=Mus musculus GN=Atp13a1	2,00E-65
BQ426348	1,00E-06	(sp Q90837) Transcriptional regulator Erg OS=Gallus gallus GN=ERG	3,00E-16
BQ426355	1,00E-06		
BQ426358	1,00E-06		
BQ426389	1,00E-06		
BQ426395	1,00E-06	predicted protein [<i>Nematostella vectensis</i>]	2,00E-19
BQ426399	1,00E-06		
BQ426407	1,00E-06		
BQ426411	1,00E-06		
BQ426428	1,00E-06		
BQ426447	1,00E-06	(sp O55135) Eukaryotic translation initiation factor 6 OS=Mus musculus GN=Eif6	1,00E-98
BQ426456	1,00E-06	(sp Q9JJI5) Neuromedin-U receptor 1 OS=Rattus norvegicus GN=Nmur1	3,00E-08
BQ426466	1,00E-06		
BQ426470	1,00E-06		
BQ426479	1,00E-06	(sp Q24799) Myophilin OS=Echinococcus granulosus	2,00E-52
BQ426481	1,00E-06	(sp Q90YV5) 60S ribosomal protein L13 OS=Ictalurus punctatus GN=rpl13	2,00E-70
BQ426484	1,00E-06		
BQ426485	1,00E-05		
BQ426497	1,00E-06		
BQ426509	1,00E-05		

BQ426511	1,00E-06	(sp Q6Q899) Probable ATP-dependent RNA helicase DDX58 OS=Mus musculus GN=Ddx58	2,00E-28
BQ426520	1,00E-06	(sp Q5SPP5) Protein EFR3 homolog B OS=Danio rerio GN=efr3b	1,00E-17
BQ426525	1,00E-05		
BQ426550	1,00E-06	(sp P12812) Major egg antigen OS=Schistosoma mansoni	7,00E-06
BQ426552	1,00E-06		
BQ426559	1,00E-06		
BQ426571	1,00E-06	(sp P62845) 40S ribosomal protein S15 OS=Rattus norvegicus GN=Rps15	2,00E-64
BQ426589	1,00E-06		
BQ426612	1,00E-06		
BQ426614	1,00E-05	PREDICTED: hypothetical protein [Ciona intestinalis]	4,00E-06
BQ426626	1,00E-06		
BQ426652	1,00E-05	(sp Q8C3L1) Uncharacterized protein C3orf32 homolog OS=Mus musculus	1,00E-41
BQ426658	1,00E-06	predicted protein [Nematostella vectensis]	3,00E-16
BQ426659	1,00E-06	(sp Q96GP6) Scavenger receptor class F member 2 OS=Homo sapiens GN=SCARF2	5,00E-07
BQ426661	1,00E-06	(sp Q69BK3) Cytochrome b-c1 complex subunit Rieske, mitochondrial OS=Pongo pygmaeus GN=UQCRFS1	1,00E-51
BQ426670	1,00E-06		
BQ426671	1,00E-06	(sp P49230) Profilin OS=Entamoeba histolytica	5,00E-08
BQ426672	1,00E-06	(sp P48601) 26S protease regulatory subunit 4 OS=Drosophila melanogaster GN=Pros26.4	0,00E+00
BQ426673	1,00E-06		
BQ426694	1,00E-05		
BQ426708	1,00E-06		
BQ426717	1,00E-06	conserved hypothetical protein [Pediculus humanus corporis]	5,00E-12
BQ426725	1,00E-06		
BQ426736	1,00E-06	hypothetical protein BRAFLDRAFT_124513 [Branchiostoma floridae]	7,00E-14
BQ426743	1,00E-06		
BQ426746	1,00E-06		
BQ426751	1,00E-06	(sp O46160) 60S ribosomal protein L14 OS=Lumbricus rubellus GN=RPL14	5,00E-46
BQ426758	1,00E-06		

BQ426761	1,00E-06	(sp Q9QY73) Transmembrane protein 59 OS=Mus musculus GN=Tmem59	5,00E-21
BQ426763	1,00E-05		
BQ426765	1,00E-06	(sp Q14974) Importin subunit beta-1 OS=Homo sapiens GN=KPNB1	3,00E-95
BQ426779	1,00E-06		
BQ426786	1,00E-06		
BQ426800	1,00E-06		
BQ426814	1,00E-06	(sp P61127) 60S ribosomal protein L32 OS=Drosophila yakuba GN=RpL32	5,00E-51
BQ426815	1,00E-06		
BQ426816	1,00E-06		
BQ426821	1,00E-06		
BQ426825	1,00E-06	(sp Q14376) UDP-glucose 4-epimerase OS=Homo sapiens GN=GALE	2,00E-12
BQ426831	1,00E-06	(sp P62329) Thymosin beta-4 OS=Rattus norvegicus GN=Tmsb4x	1,00E-12
BQ426851	1,00E-06		
BQ426873	1,00E-06	(sp P61980) Heterogeneous nuclear ribonucleoprotein K OS=Rattus norvegicus GN=Hnrnpk	1,00E-18
BQ426885	1,00E-06		
BQ426891	1,00E-06		
BQ426894	1,00E-06	(sp Q9EQ06) Estradiol 17-beta-dehydrogenase 11 OS=Mus musculus GN=Hsd17b11	1,00E-53
BQ426904	1,00E-06		
BQ426908	1,00E-06		
BQ426925	1,00E-06		
BQ426932	1,00E-06	(sp P83686) NADH-cytochrome b5 reductase 3 (Fragment) OS=Sus scrofa GN=CYB5R3	1,00E-108
BQ426945	1,00E-06		
BQ426950	1,00E-06		
BQ426954	1,00E-05	(sp Q8C3J5) Deducator of cytokinesis protein 2 OS=Mus musculus GN=Dock2	6,00E-34
BQ426957	1,00E-05	(sp P37889) Fibulin-2 OS=Mus musculus GN=Fbln2	5,00E-28
BQ426959	1,00E-06		
BQ426962	1,00E-06		
BQ426997	1,00E-06	(sp P68372) Tubulin beta-2C chain OS=Mus musculus GN=Tubb2c	0,00E+00
BQ426999	1,00E-06	(sp Q9D1A0) UPF0308 protein C9orf21 homolog OS=Mus musculus	5,00E-07
BQ427003	1,00E-06		

BQ427007	1,00E-06		
BQ427020	1,00E-06		
BQ427025	1,00E-06		
BQ427028	1,00E-06	(sp Q5E946) Protein DJ-1 OS=Bos taurus GN=PARK7	1,00E-51
BQ427029	1,00E-06	(sp P06603) Tubulin alpha-1 chain OS=Drosophila melanogaster	
BQ427040	1,00E-06	GN=alphaTub84B	0,00E+00
BQ427052	1,00E-06	(sp P62972) Ubiquitin OS=Xenopus laevis	5,00E-34
BQ427073	1,00E-06		
BQ427098	1,00E-06		
BQ427110	1,00E-06	(sp Q5R9R4) 60S ribosomal protein L7 OS=Pongo abelii GN=RPL7	2,00E-89
BQ427140	1,00E-06		
BQ427144	1,00E-06	(sp Q7TNC8) Glycine receptor subunit alpha-2 OS=Mus musculus GN=Glr2	4,00E-41
BQ427153	1,00E-06		
BQ427200	1,00E-06		
BQ427204	1,00E-06		
BQ427206	1,00E-06		
BQ427208	1,00E-06		
BQ427216	1,00E-06		
BQ427225	1,00E-06	(sp POC2C1) 39S ribosomal protein L30, mitochondrial OS=Rattus norvegicus	
BQ427228	1,00E-06	GN=Mrpl30	1,00E-11
BQ427237	1,00E-06		
BQ427253	1,00E-06	(sp Q5ZME2) Malate dehydrogenase, cytoplasmic OS=Gallus gallus	
BQ427254	1,00E-05	GN=MDH1	3,00E-41
BQ427294	1,00E-06	(sp P27076) 60S ribosomal protein L44 OS=Kluyveromyces marxianus	
BQ427295	1,00E-05	GN=RPL44	1,00E-33
BQ427310	1,00E-05	(sp Q9NB34) 60S ribosomal protein L34 OS=Aedes triseriatus GN=RpL34	7,00E-39
BQ427312	1,00E-06	(sp P08645) Ras-like protein 3 OS=Drosophila melanogaster GN=R	6,00E-75
BQ427314	1,00E-06	(sp Q86WV5) Uncharacterized protein FLJ39785 OS=Homo sapiens	1,00E-09
BQ427318	1,00E-06	(sp Q96KG7) Multiple epidermal growth factor-like domains 10 OS=Homo sapiens GN=MEGF10	8,00E-06
		(sp Q14247) Src substrate cortactin OS=Homo sapiens GN=CTTN	1,00E-123

BQ427330	1,00E-06		
BQ427337	1,00E-06		
BQ427342	1,00E-05		
BQ427347	1,00E-06		
BQ427355	1,00E-06	(sp P52735) Guanine nucleotide exchange factor VAV2 OS=Homo sapiens GN=VAV2	1,00E-39
CK172314	1,00E-06	(sp P02637) Sarcoplasmic calcium-binding protein OS=Patinopecten yessoensis	2,00E-55
CK172322	1,00E-06		
CK172325	1,00E-06		
CK172328	1,00E-06		
CK172339	1,00E-06		
CK172365	1,00E-05	(sp P00399) Cytochrome c oxidase subunit 1 OS=Drosophila melanogaster GN=mt:CoI	1,00E-137
CK172373	1,00E-06		
CK172374	1,00E-06	Chlamydomonas reinhardtii strain CC-503 cw92 mt+	2,00E-09
CK172383	1,00E-06	(sp Q61M78) DNA topoisomerase I, mitochondrial OS=Rattus norvegicus GN=Top1mt	2,00E-09
CK172392	1,00E-06		
CK172394	1,00E-06		
CU681473	1,00E-06		
CU681477	1,00E-05	(sp P28288) ATP-binding cassette sub-family D member 3 OS=Homo sapiens GN=ABCD3	1,00E-151
CU681483	1,00E-06	(sp P00179) Cytochrome P450 2C5 OS=Oryctolagus cuniculus GN=CYP2C5	3,00E-34
CU681485	1,00E-06		
CU681491	1,00E-06		
CU681495	1,00E-06	(sp Q7SY29) Sugar phosphate exchanger 2 OS=Danio rerio GN=slc37a2	2,00E-72
CU681496	1,00E-06		
CU681497	1,00E-06		
CU681499	1,00E-06	PREDICTED: similar to palmitoyl-protein thioesterase [Acyrtosiphon pisum] (sp Q28943) Dihydropyrimidine dehydrogenase [NADP+] OS=Sus scrofa GN=DPYD	4,00E-06
CU681507	1,00E-06		9,00E-36
CU681508	1,00E-05		

CU681511	1,00E-06	(sp Q9VZZ4) Peroxidase OS=Drosophila melanogaster GN=Pxn	2,00E-07
CU681518	1,00E-06	PREDICTED: similar to predicted protein [Hydra magnipapillata]	3,00E-06
CU681526	1,00E-05	(sp Q5R9L3) Ras GTPase-activating protein-binding protein 2 OS=Pongo abelii GN=G3BP2	7,00E-37
CU681528	1,00E-06	(sp Q5M7G4) Putative adenylate kinase-like protein C9orf98 homolog OS=Xenopus laevis	6,00E-32
CU681529	1,00E-06	(sp P55780) NADH-ubiquinone oxidoreductase chain 2 OS=Gadus morhua GN=MT-ND2	5,00E-11
CU681533	1,00E-06		
CU681538	1,00E-06		
CU681540	1,00E-06		
CU681545	1,00E-05	(sp Q5R4J9) NMDA receptor-regulated protein 1 OS=Pongo abelii GN=NARG1	2,00E-22
CU681553	1,00E-05		
CU681561	1,00E-06	(sp Q5M8V0) Transcription factor BTF3 homolog 4 OS=Xenopus tropicalis GN=btf3l4	2,00E-52
CU681564	1,00E-06		
CU681569	1,00E-06		
CU681573	1,00E-06		
CU681577	1,00E-06	(sp P30405) Peptidyl-prolyl cis-trans isomerase, mitochondrial OS=Homo sapiens GN=PPIF	1,00E-72
CU681582	1,00E-06	hypothetical protein BRAFLDRAFT_74086 [Branchiostoma floridae]	2,00E-10
CU681583	1,00E-06		
CU681587	1,00E-06	(sp P55043) GTP-binding protein RAD OS=Rattus norvegicus GN=Rrad	5,00E-13
CU681593	1,00E-06	(sp Q0MQ86) NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 12 OS=Gorilla gorilla gorilla	1,00E-27
CU681594	1,00E-06	(sp P35486) Pyruvate dehydrogenase E1 component subunit alpha, somatic form, mitochondrial OS=Mus musculus	1,00E-131
CU681598	1,00E-06		
CU681599	1,00E-06		
CU681603	1,00E-06	(sp Q9H6D7) HAUS augmin-like complex subunit 4 OS=Homo sapiens GN=HAUS4	1,00E-23
CU681616	1,00E-06	(sp O88413) Tubby-related protein 3 OS=Mus musculus GN=Tulp3	6,00E-32
CU681618	1,00E-06		

CU681620	1,00E-06	(sp P04179) Superoxide dismutase [Mn], mitochondrial OS=Homo sapiens GN=SOD2	4,00E-75
CU681624	1,00E-06	(sp P49614) Beta-hexosaminidase subunit beta OS=Felis catus GN=HEXB	1,00E-151
CU681630	1,00E-06		
CU681638	1,00E-06	hypothetical protein BRAFLDRAFT_83240 [Branchiostoma floridae]	1,00E-06
CU681640	1,00E-06	(sp Q9VUY9) Phosphoglucomutase OS=Drosophila melanogaster GN=Pgm	1,00E-114
CU681658	1,00E-06		
CU681659	1,00E-05		
CU681669	1,00E-06	(sp Q13201) Multimerin-1 OS=Homo sapiens GN=MMRN1	3,00E-06
CU681670	1,00E-06	(sp Q7Z0T3) Temptin OS=Aplysia californica	4,00E-22
CU681677	1,00E-05	(sp Q13620) Cullin-4B OS=Homo sapiens GN=CUL4B	3,00E-14
CU681679	1,00E-06	(sp Q10751) Angiotensin-converting enzyme (Fragment) OS=Gallus gallus GN=ACE	7,00E-37
CU681681	1,00E-06	(sp Q1JPZ7) Pre-mRNA-processing factor 39 OS=Danio rerio GN=prpf39	2,00E-14
CU681683	1,00E-05	(sp Q6GMG8) Lipopolysaccharide-induced tumor necrosis factor-alpha factor homolog OS=Danio rerio	7,00E-18
CU681691	1,00E-06	(sp Q9DG68) 60S acidic ribosomal protein P0 OS=Rana sylvatica GN=RPLP0	1,00E-105
CU681695	1,00E-06		
CU681709	1,00E-06	(sp Q6DDU9) RNA-binding protein 5-B OS=Xenopus laevis GN=rbm5-B	2,00E-23
CU681713	1,00E-06	(sp O35474) EGF-like repeat and discoidin I-like domain-containing protein 3 OS=Mus musculus	5,00E-12
CU681716	1,00E-06	(sp Q5ZM98) Stress-70 protein, mitochondrial OS=Gallus gallus GN=HSPA9	0,00E+00
CU681727	1,00E-06	(sp Q62203) Splicing factor 3A subunit 2 OS=Mus musculus GN=Sf3a2	6,00E-81
CU681730	1,00E-06		
CU681740	1,00E-05		
CU681750	1,00E-05		
CU681755	1,00E-06	(sp Q6NYL3) Peroxisomal bifunctional enzyme OS=Danio rerio GN=ehhadh	4,00E-40
CU681759	1,00E-06		
CU681762	1,00E-06	PREDICTED: similar to superoxide dismutase [Strongylocentrotus purpuratus]	3,00E-06
CU681764	1,00E-06	expressed hypothetical protein [Trichoplax adhaerens]	1,00E-12

CU681768	1,00E-06		
CU681779	1,00E-06		
CU681784	1,00E-06	(sp P15253) Calreticulin OS= <i>Oryctolagus cuniculus</i> GN=CALR	1,00E-124
CU681793	1,00E-05	(sp Q6DIB5) Multiple epidermal growth factor-like domains 10 OS= <i>Mus musculus</i> GN=Megf10	8,00E-08
CU681795	1,00E-06	(sp O75897) Sulfotransferase 1C4 OS= <i>Homo sapiens</i> GN=SULT1C4	4,00E-30
CU681798	1,00E-06		
CU681812	1,00E-06	(sp Q921M3) Splicing factor 3B subunit 3 OS= <i>Mus musculus</i> GN=Sf3b3	5,00E-18
CU681815	1,00E-06		
CU681816	1,00E-06	(sp Q8HXX6) Ganglioside GM2 activator OS= <i>Macaca fascicularis</i> GN=GM2A	2,00E-22
CU682012	1,00E-06		
CU682018	1,00E-06	PREDICTED: similar to superoxide dismutase [<i>Strongylocentrotus purpuratus</i>]	3,00E-06
CU682024	1,00E-06		
CU682063	1,00E-06	(sp Q61191) Host cell factor OS= <i>Mus musculus</i> GN=Hcfc1	1,00E-82
CU682079	1,00E-05	(sp P49793) Nuclear pore complex protein Nup98 OS= <i>Rattus norvegicus</i> GN=Nup98	3,00E-55
CU682081	1,00E-06	hypothetical protein LOC734261 [<i>Xenopus laevis</i>]	2,00E-06
CU682086	1,00E-05		
CU682087	1,00E-06	(sp A6H584) Collagen alpha-5(VI) chain OS= <i>Mus musculus</i> GN=Col29a1	2,00E-17
CU682090	1,00E-06		
CU682094	1,00E-06	(sp Q6NVP6) U4/U6 small nuclear ribonucleoprotein Prp31 OS= <i>Xenopus tropicalis</i> GN=prpf31	8,00E-54
CU682095	1,00E-06		
CU682117	1,00E-06	(sp Q9NY25) C-type lectin domain family 5 member A OS= <i>Homo sapiens</i> GN=CLEC5A	3,00E-06
CU682119	1,00E-06	(sp O60294) Leucine carboxyl methyltransferase 2 OS= <i>Homo sapiens</i> GN=LCMT2	2,00E-10
CU682121	1,00E-06		
CU682122	1,00E-06		
CU682126	1,00E-05		
CU682137	1,00E-06	(sp Q20FR3) Cytochrome b OS= <i>Lepilemur edwardsi</i> GN=MT-CYB	5,00E-41
CU682142	1,00E-06	(sp Q7Z0T3) Temptin OS= <i>Aplysia californica</i>	4,00E-20

CU682156	1,00E-06	(sp Q5R407) Ubiquitin carboxyl-terminal hydrolase 5 OS=Pongo abelii GN=UBP5	4,00E-21
CU682166	1,00E-05		
CU682176	1,00E-06		
CU682180	1,00E-06		
CU682183	1,00E-06	PREDICTED: similar to Os09g0542200 [Strongylocentrotus purpuratus]	1,00E-12
CU682184	1,00E-06	hypothetical protein BRAFLDRAFT_208032 [Branchiostoma floridae]	5,00E-25
CU682186	1,00E-06		
CU682192	1,00E-06	(sp P34713) Multidrug resistance protein 3 OS=Caenorhabditis elegans GN=pgp-3	4,00E-66
CU682195	1,00E-05		
CU682197	1,00E-06	(sp P18420) Proteasome subunit alpha type-1 OS=Rattus norvegicus GN=Psm1	1,00E-55
CU682198	1,00E-06	(sp Q9JI20) Double-stranded RNA-specific editase B2 OS=Mus musculus GN=Adarb2	3,00E-06
CU682207	1,00E-05	(sp Q3UGF1) WD repeat-containing protein 19 OS=Mus musculus GN=Wdr19	5,00E-26
CU682210	1,00E-05		
CU682215	1,00E-06	hypothetical protein BRAFLDRAFT_88895 [Branchiostoma floridae]	1,00E-12
CU682229	1,00E-06		
CU682231	1,00E-06		
CU682242	1,00E-06		
CU682260	1,00E-06		
CU682272	1,00E-06	hypothetical protein BRAFLDRAFT_119590 [Branchiostoma floridae]	5,00E-47
CU682273	1,00E-06	(sp P37892) Carboxypeptidase E OS=Lophius americanus GN=cpe	3,00E-19
CU682291	1,00E-06		
CU682295	1,00E-06		
CU682301	1,00E-06	(sp P11954) Threonine dehydratase catabolic OS=Salmonella typhimurium GN=tdcB	9,00E-40
CU682313	1,00E-05		
CU682314	1,00E-05		
CU682324	1,00E-06	(sp Q3SZ90) 60S ribosomal protein L13a OS=Bos taurus GN=RPL13A	7,00E-71
CU682332	1,00E-06		
CU682333	1,00E-06		
CU682385	1,00E-06		

CU682395	1,00E-06	hypothetical protein BRAFLDRAFT_123768 [Branchiostoma floridae]	1,00E-09
CU682405	1,00E-06		
CU682406	1,00E-05		
CU682408	1,00E-06	(sp A6QLA6) Chromatin assembly factor 1 subunit A OS=Bos taurus	
CU682413	1,00E-06	GN=CHAF1A	3,00E-23
CU682418	1,00E-06	(sp P12527) Arachidonate 5-lipoxygenase OS=Rattus norvegicus GN=Alox5	3,00E-39
CU682430	1,00E-06	(sp Q8STF0) Calmodulin OS=Strongylocentrotus intermedius	3,00E-79
CU682440	1,00E-06	(sp Q17Q91) Coiled-coil-helix-coiled-coil-helix domain-containing protein 7	
CU682450	1,00E-05	OS=Bos taurus GN=CHCHD7	2,00E-09
CU682471	1,00E-06	(sp O93574) Reelin (Fragment) OS=Gallus gallus GN=RELN	1,00E-29
CU682485	1,00E-06	(sp Q81TC3) 40S ribosomal protein S19 OS=Aequipecten irradians GN=RPS19	2,00E-58
CU682489	1,00E-06		
CU682507	1,00E-06	(sp Q6QLW4) Cytochrome c OS=Pectinaria gouldii	9,00E-50
CU682521	1,00E-06		
CU682527	1,00E-05		
CU682536	1,00E-06	(sp Q7Z0T3) Temptin OS=Aplysia californica	1,00E-08
CU682553	1,00E-06		
CU682571	1,00E-06		
CU682574	1,00E-06	Xenopus laevis hypothetical protein LOC100049085 (LOC100049085), mRNA	2,00E-06
CU682577	1,00E-06	(sp Q32NR4) Tetratricopeptide repeat protein 29 OS=Xenopus laevis	
CU682583	1,00E-06	GN=ttc29	1,00E-34
CU682602	1,00E-06		
CU682609	1,00E-05		
CU682613	1,00E-05	(sp Q7SZ59) Ras-like protein family member 12 OS=Danio rerio GN=RASL12	6,00E-40
CU682631	1,00E-06	(sp Q99P88) Nuclear pore complex protein Nup155 OS=Mus musculus	
CU682672	1,00E-06	GN=Nup155	2,00E-25
		(sp Q5RE69) Nepriylisin OS=Pongo abelii GN=MME	4,00E-67

CU682700	1,00E-05	(sp Q9W265) Probable hydroxyacid-oxoacid transhydrogenase, mitochondrial OS=Drosophila melanogaster	3,00E-22
CU682713	1,00E-06		
CU682728	1,00E-06		
CU682732	1,00E-06		
CU682754	1,00E-06	(sp Q7M370) Arylacetamide deacetylase OS=Oryctolagus cuniculus GN=AADAC	3,00E-11
CU682770	1,00E-06		
CU682787	1,00E-06		
CU682794	1,00E-06	(sp P12111) Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 SV=4	8,00E-07
CU682798	1,00E-06		
CU682816	1,00E-06	(sp Q3TIX9) U4/U6.U5 tri-snRNP-associated protein 2 OS=Mus musculus GN=Usp39	1,00E-137
CU682818	1,00E-06		
CU682833	1,00E-06		
CU682834	1,00E-06		
CU682846	1,00E-05	(sp Q9C040) Tripartite motif-containing protein 2 OS=Homo sapiens GN=TRIM2	8,00E-18
CU682851	1,00E-06	(sp P23342) Monomeric sarcosine oxidase OS=Bacillus sp. (strain NS-129) GN=soxA	5,00E-45
CU682853	1,00E-06		
CU682860	1,00E-05	(sp P07871) 3-ketoacyl-CoA thiolase B, peroxisomal OS=Rattus norvegicus GN=Acaa1b	8,00E-50
CU682868	1,00E-06		
CU682887	1,00E-06	(sp O95340) Bifunctional 3'-phosphoadenosine 5'-phosphosulfate synthetase 2 OS=Homo sapiens	3,00E-16
CU682896	1,00E-06	(sp Q9P278) Folliculin-interacting protein 2 OS=Homo sapiens GN=FNIP2 PREDICTED: similar to spermatogenesis associated 4 [Strongylocentrotus purpuratus]	2,00E-09
CU682905	1,00E-06		1,00E-07
CU682906	1,00E-06		
CU682908	1,00E-06	(sp Q27HV0) UDP-N-acetylglucosamine--peptide N-acetylglucosaminyltransferase 110 kDa subunit OS=Sus scrofa	2,00E-68
CU682915	1,00E-06	(sp Q28062) Brevican core protein OS=Bos taurus GN=BCAN	2,00E-06

CU682919	1,00E-06	(sp Q5E9E4) Translocon-associated protein subunit beta OS=Bos taurus GN=SSR2	6,00E-62
CU682926	1,00E-06	(sp O49923) Adenosine kinase OS=Physcomitrella patens GN=ADK	1,00E-65
CU682934	1,00E-06		
CU682939	1,00E-06		
CU682940	1,00E-06		
CU682946	1,00E-06	hypothetical protein BRAFLDRAFT_89746 [Branchiostoma floridae] (sp O15244) Solute carrier family 22 member 2 OS=Homo sapiens GN=SLC22A2	8,00E-07 9,00E-25
CU682969	1,00E-06		
CU682971	1,00E-06		
CU682973	1,00E-06		
CU682991	1,00E-06		
CU682993	1,00E-06		
CU682996	1,00E-06	(sp Q30HU9) Serine protease inhibitor Cvs1-1 OS=Crassostrea virginica	1,00E-06
CU683009	1,00E-06		
CU683023	1,00E-06		
CU683025	1,00E-06		
CU683034	1,00E-06	(sp P49900) Arginase, hepatic OS=Rana catesbeiana	4,00E-90
CU683037	1,00E-06	(sp O76840) Papilin OS=Caenorhabditis elegans GN=mig-6	8,00E-10
CU683038	1,00E-06		
CU683043	1,00E-06	(sp Q5Z187) Tubulin-specific chaperone D OS=Gallus gallus GN=TBCD	7,00E-22
CU683056	1,00E-06		
CU683075	1,00E-06		
CU683077	1,00E-06	(sp O02649) 60 kDa heat shock protein, mitochondrial OS=Drosophila melanogaster GN=Hsp60	3,00E-76
CU683103	1,00E-05	(sp Q5F384) Protein YIPF3 OS=Gallus gallus GN=YIPF3	2,00E-28
CU683105	1,00E-05	(sp B5X171) Pescadillo homolog OS=Salmo salar GN=pes1	2,00E-53
CU683107	1,00E-06	(sp P43245) Multidrug resistance protein 1 OS=Rattus norvegicus GN=Abcb1 (sp Q27128) Bifunctional 3'-phosphoadenosine 5'-phosphosulfate synthetase	1,00E-34
CU683112	1,00E-06	OS=Urechis caupo	7,00E-54
CU683115	1,00E-06	(sp Q5EB28) Midnolin OS=Xenopus tropicalis GN=midn	1,00E-07
CU683120	1,00E-05		
CU683127	1,00E-06		
CU683151	1,00E-06		

CU683160	1,00E-06	(sp Q8BR76) Meckelin OS=Mus musculus GN=Tmem67	4,00E-33
CU683161	1,00E-06	(sp A2ASS6) Titin OS=Mus musculus GN=Ttn	6,00E-23
CU683171	1,00E-05		
CU683178	1,00E-06	(sp Q5T2S8) Armadillo repeat-containing protein 4 OS=Homo sapiens GN=ARMC4	1,00E-21
CU683196	1,00E-06		
CU683208	1,00E-06	(sp Q12841) Follistatin-related protein 1 OS=Homo sapiens GN=FSTL1 (sp Q9GP32) Fructose-bisphosphate aldolase OS=Echinococcus multilocularis GN=FBPA	5,00E-28
CU683220	1,00E-06	(sp Q5RE52) Anaphase-promoting complex subunit 5 OS=Pongo abelii GN=ANAPC5	2,00E-28
CU683221	1,00E-06		4,00E-10
CU683225	1,00E-06	(sp P00789) Calpain-1 catalytic subunit OS=Gallus gallus	1,00E-28
CU683227	1,00E-06		
CU683233	1,00E-06	(sp Q08AE8) Protein spire homolog 1 OS=Homo sapiens GN=SPIRE1	4,00E-32
CU683236	1,00E-06		
CU683259	1,00E-06	(sp Q4FZY0) EF-hand domain-containing protein D2 OS=Rattus norvegicus GN=Efhd2	5,00E-48
CU683261	1,00E-06	(sp O95428) Papilin OS=Homo sapiens GN=PAPLN SV=4 (sp Q9H6T0) Epithelial splicing regulatory protein 2 OS=Homo sapiens GN=ESRP2	3,00E-27
CU683266	1,00E-05		1,00E-19
CU683271	1,00E-06		
CU683277	1,00E-06		
CU683285	1,00E-06	(sp Q99KY4) Cyclin G-associated kinase OS=Mus musculus GN=Gak	5,00E-83
CU683294	1,00E-06		
CU683296	1,00E-06	(sp Q62415) Apoptosis-stimulating of p53 protein 1 OS=Mus musculus GN=Ppp1r13b	7,00E-12
CU683298	1,00E-06		
CU683307	1,00E-06		
CU683310	1,00E-06		
CU683312	1,00E-06		
CU683315	1,00E-06	(sp P11217) Glycogen phosphorylase, muscle form OS=Homo sapiens GN=PYGM SV=6	0,00E+00
CU683317	1,00E-05	(sp Q6GQT0) Dual specificity protein phosphatase CDC14A OS=Mus musculus GN=Cdc14a	3,00E-06
CU683321	1,00E-05		

CU683323	1,00E-06	(sp Q28042) Oviduct-specific glycoprotein (Fragment) OS=Bos taurus GN=OVGP1	1,00E-49
CU683331	1,00E-05	(sp P42893) Endothelin-converting enzyme 1 OS=Rattus norvegicus GN=Ece1	2,00E-55
CU683332	1,00E-06		
CU683333	1,00E-06	(sp P21589) 5'-nucleotidase OS=Homo sapiens GN=NT5E	2,00E-10
CU683337	1,00E-06	(sp A9C3P0) Malectin OS=Danio rerio GN=mlec	5,00E-17
CU683341	1,00E-05		
CU683345	1,00E-06	(sp Q6AXS9) Protein FAM101B OS=Rattus norvegicus GN=Fam101b	9,00E-09
CU683353	1,00E-06		
CU683363	1,00E-06	(sp Q99675) Cell growth regulator with RING finger domain protein 1 OS=Homo sapiens GN=CGRRF1	5,00E-23
CU683365	1,00E-06		
CU683371	1,00E-06		
CU683376	1,00E-06	(sp P62994) Growth factor receptor-bound protein 2 OS=Rattus norvegicus GN=Grb2	4,00E-70
CU683379	1,00E-06	(sp P43427) Solute carrier family 2, facilitated glucose transporter member 5 OS=Rattus norvegicus GN=Slc2a5	6,00E-14
CU683387	1,00E-06		
CU683397	1,00E-06		
CU683414	1,00E-06	(sp Q64FW2) All-trans-retinol 13,14-reductase OS=Mus musculus GN=Retsat	7,00E-75
CU683415	1,00E-06	(sp Q8K0V4) CCR4-NOT transcription complex subunit 3 OS=Mus musculus GN=Cnot3	7,00E-86
CU683421	1,00E-06		
CU683427	1,00E-06		
CU683446	1,00E-06	(sp O95428) Papilin OS=Homo sapiens GN=PAPLN SV=4	3,00E-27
CU683464	1,00E-06		
CU683466	1,00E-06		
CU683474	1,00E-06		
CU683476	1,00E-06	PREDICTED: hypothetical protein [Rattus norvegicus]	5,00E-06
CU683480	1,00E-06		
CU683483	1,00E-05	(sp Q62504) Msx2-interacting protein OS=Mus musculus GN=Spen	2,00E-20
CU683488	1,00E-06		
CU683491	1,00E-06		

CU683497	1,00E-06	(sp Q54MV2) Probable serine/threonine-protein kinase MARK-B	
CU683513	1,00E-06	OS=Dictyostelium discoideum GN=mrkB	3,00E-13
CU683516	1,00E-06	(sp Q9Y614) Ubiquitin carboxyl-terminal hydrolase 3	
CU683536	1,00E-06	GN=USP3 OS=Homo sapiens	8,00E-61
CU683544	1,00E-06		
CU683547	1,00E-06	(sp P26640) Valyl-tRNA synthetase	
CU683550	1,00E-06	OS=Homo sapiens GN=VARS SV=4	5,00E-25
CU683559	1,00E-06	(sp Q81Y18) Structural maintenance of chromosomes protein 5	
CU683563	1,00E-06	sapiens GN=SMC5	5,00E-21
CU683564	1,00E-06	(sp Q9GZQ4) Neuromedin-U receptor 2	
CU683571	1,00E-05	OS=Homo sapiens GN=NMUR2	2,00E-11
CU683572	1,00E-05	(sp Q15643) Thyroid receptor-interacting protein 11	
CU683574	1,00E-06	OS=Homo sapiens GN=TRIP11	8,00E-31
CU683576	1,00E-06	(sp Q29116) Tenascin	
CU683577	1,00E-06	OS=Sus scrofa GN=TNC	1,00E-06
CU683585	1,00E-06	(sp Q8HZK2) Dual oxidase 2	
CU683587	1,00E-06	OS=Sus scrofa GN=DUOX2	3,00E-26
CU683609	1,00E-06	(sp Q5R7M2) D-3-phosphoglycerate dehydrogenase	
CU683610	1,00E-06	OS=Pongo abelii GN=PHGDH	4,00E-26
CU683614	1,00E-06	(sp Q8UWA5) Carbonic anhydrase 2	
CU683624	1,00E-06	OS=Tribolodon hakonensis GN=ca2	3,00E-24
CU683628	1,00E-06	(sp O99824) NADH-ubiquinone oxidoreductase chain 1	
CU683646	1,00E-05	OS=Rhipicephalus sanguineus GN=ND1	9,00E-42
CU683647	1,00E-06		
CU683659	1,00E-06	(sp P18484) AP-2 complex subunit alpha-2	
CU683661	1,00E-06	OS=Rattus norvegicus GN=Ap2a2	8,00E-78
CU683662	1,00E-06	(sp P13395) Spectrin alpha chain	
		OS=Drosophila melanogaster GN=alpha-Spec	1,00E-74

CU683672	1,00E-06		
CU683674	1,00E-05	predicted protein [<i>Nematostella vectensis</i>]	9,00E-08
CU683675	1,00E-06	(sp Q5R4R7) Protein KIAA1045 homolog OS= <i>Pongo abelii</i>	8,00E-30
CU683682	1,00E-06		
CU683687	1,00E-06		
CU683688	1,00E-05	(sp Q9NZR2) Low-density lipoprotein receptor-related protein 1B OS= <i>Homo sapiens</i> GN=LRP1B	5,00E-11
CU683695	1,00E-06	(sp P31639) Sodium/glucose cotransporter 2 OS= <i>Homo sapiens</i> GN=SLC5A2	9,00E-26
CU683698	1,00E-06	(sp Q9VZF4) F-box/WD repeat-containing protein 7 OS= <i>Drosophila melanogaster</i> GN=ago	4,00E-13
CU683710	1,00E-05	(sp Q9VXK0) Protein NipSnap OS= <i>Drosophila melanogaster</i> GN=Nipsnap	7,00E-67
CU683720	1,00E-05		
CU683721	1,00E-06	(sp Q8NFD2) Ankyrin repeat and protein kinase domain-containing protein 1 OS= <i>Homo sapiens</i> GN=ANKK1	3,00E-10
CU683728	1,00E-06		
CU683733	1,00E-06		
CU683777	1,00E-06		
CU683784	1,00E-06	(sp Q37374) Cytochrome c oxidase subunit 3 OS= <i>Acanthamoeba castellanii</i> GN=COX3	2,00E-18
CU683786	1,00E-06		
CU684059	1,00E-06		
CU684068	1,00E-06	(sp P19804) Nucleoside diphosphate kinase B OS= <i>Rattus norvegicus</i> GN=Nme2	2,00E-58
CU684071	1,00E-06		
CU684085	1,00E-05	(sp Q6UWW8) Carboxylesterase 3 OS= <i>Homo sapiens</i> GN=CES3	6,00E-52
CU684094	1,00E-06	(sp Q6ZQ08) CCR4-NOT transcription complex subunit 1 OS= <i>Mus musculus</i> GN=Cnot1	1,00E-146
CU684105	1,00E-06		
CU684106	1,00E-06	(sp Q12872) Splicing factor, arginine/serine-rich 8 OS= <i>Homo sapiens</i> GN=SFRS8	3,00E-11
CU684108	1,00E-06		
CU684125	1,00E-06		
CU684127	1,00E-05	(sp Q61425) Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS= <i>Mus musculus</i> GN=Hadh	5,00E-47

CU684172	1,00E-06	(sp Q11011) Puromycin-sensitive aminopeptidase OS=Mus musculus GN=Npepps	3,00E-81
CU684178	1,00E-06	(sp Q9N0N3) Transcription factor AP-2 alpha OS=Ovis aries GN=TFAP2A	7,00E-44
CU684184	1,00E-06		
CU684211	1,00E-06		
CU684214	1,00E-06		
CU684221	1,00E-06		
CU684236	1,00E-06	(sp Q0P467) Ribonuclease kappa-A OS=Danio rerio GN=rnaseka	5,00E-21
CU684251	1,00E-06		
CU684258	1,00E-06	(sp Q8JGT4) Netrin receptor UNC5B OS=Xenopus laevis GN=unc5b	4,00E-09
CU684265	1,00E-06		
CU684269	1,00E-06		
CU684280	1,00E-06	hypothetical protein [Schistosoma mansoni]	8,00E-13
CU684298	1,00E-05	PREDICTED: hypothetical protein [Strongylocentrotus purpuratus]	5,00E-06
		(sp Q5ZL33) Serine-threonine kinase receptor-associated protein OS=Gallus gallus GN=STRAP	1,00E-118
CU684314	1,00E-06		
CU684315	1,00E-06		
CU684316	1,00E-06		
		(sp P20478) Glutamine synthetase 2 cytoplasmic OS=Drosophila melanogaster GN=Gs2	2,00E-09
CU684320	1,00E-06		
		(sp P82198) Transforming growth factor-beta-induced protein ig-h3 OS=Mus musculus GN=Tgfb1	2,00E-12
CU684323	1,00E-06		
CU684327	1,00E-06		
		(sp P00399) Cytochrome c oxidase subunit 1 OS=Drosophila melanogaster GN=mt:Co1	1,00E-137
CU684332	1,00E-06		
CU684350	1,00E-05		
		(sp Q9UKW6) ETS-related transcription factor Elf-5 OS=Homo sapiens GN=ELF5	1,00E-19
CU684351	1,00E-05		
CU684355	1,00E-06		
CU684366	1,00E-05	(sp C7G0B5) PIF OS=Pinctada fucata	3,00E-28
		(sp Q9UBC3) DNA (cytosine-5)-methyltransferase 3B OS=Homo sapiens GN=DNMT3B	4,00E-17
CU684371	1,00E-06		
		(sp Q9V5N8) Protocadherin-like wing polarity protein stan OS=Drosophila melanogaster GN=stan SV=4	2,00E-13
CU684379	1,00E-06		
CU684381	1,00E-06		

CU684406	1,00E-06		
CU684410	1,00E-06		
CU684418	1,00E-05		
CU684428	1,00E-06		
CU684435	1,00E-05		
CU684448	1,00E-06		
CU684456	1,00E-06		
CU684498	1,00E-06		
CU684509	1,00E-06	(sp Q24537) High mobility group protein DSP1 OS=Drosophila melanogaster GN=Dsp1	4,00E-51
CU684515	1,00E-06	(sp Q00232) NADH-ubiquinone oxidoreductase chain 5 OS=Mytilus edulis GN=ND5	7,00E-50
CU684521	1,00E-06		
CU684526	1,00E-06	Plasmodium falciparum 3D7 protein phosphatase, putative (MAL8P1.108) partial mRNA	5,00E-06
CU684533	1,00E-06		
CU684546	1,00E-06	(sp Q29CT2) Homeobox protein extradenticle OS=Drosophila pseudoobscura pseudoobscura GN=exd	6,00E-59
CU684562	1,00E-06		
CU684570	1,00E-06	(sp O35593) 26S proteasome non-ATPase regulatory subunit 14 OS=Mus musculus GN=Psm14	1,00E-127
CU684593	1,00E-06		
CU684597	1,00E-06	(sp O79417) Cytochrome c oxidase subunit 2 OS=Branchiostoma lanceolatum GN=COII	2,00E-25
CU684624	1,00E-06	(sp P03851) Uncharacterized 9.4 kDa protein OS=Escherichia coli PE=4	1,00E-13
CU684625	1,00E-05	(sp P98133) Fibrillin-1 OS=Bos taurus GN=FBN1	1,00E-14
CU684636	1,00E-06		
CU684656	1,00E-06		
CU684689	1,00E-06		
CU684691	1,00E-06		
CU684713	1,00E-06	(sp Q26261) Netrin receptor unc-5 OS=Caenorhabditis elegans GN=unc-5	6,00E-07
CU684729	1,00E-06		
CU684730	1,00E-06		
CU684732	1,00E-05		
CU684735	1,00E-06	(sp Q8CDG5) UPF0474 protein C5orf41 homolog OS=Mus musculus	3,00E-40

CU684744	1,00E-06	(sp Q64419) Liver carboxylesterase OS=Mesocricetus auratus	1,00E-09
CU684757	1,00E-06		
CU684760	1,00E-05	(sp Q2TBP4) WD repeat-containing protein 51A OS=Bos taurus GN=WDR51A	8,00E-88
CU684761	1,00E-05	(sp Q985M5) Choline dehydrogenase OS=Rhizobium loti GN=betA	9,00E-12
CU684765	1,00E-06	(sp Q5S003) Sperm-associated antigen 17 OS=Mus musculus GN=Spag17	1,00E-29
CU684768	1,00E-06	(sp Q9U639) Heat shock 70 kDa protein cognate 4 OS=Manduca sexta	0,00E+00
CU684790	1,00E-06		
CU684808	1,00E-06		
CU684809	1,00E-06	(sp Q3BCR8) Thiopurine S-methyltransferase OS=Pan troglodytes GN=TPMT	1,00E-48
CU684816	1,00E-06	(sp P17710) Hexokinase-1 OS=Mus musculus GN=HK1	1,00E-116
CU684836	1,00E-06	(sp Q5BIM8) DNA excision repair protein ERCC-8 OS=Bos taurus GN=ERCC8	1,00E-17
CU684842	1,00E-06		
CU684844	1,00E-05		
CU684862	1,00E-05		
CU684864	1,00E-06	(sp Q8NE09) Regulator of G-protein signaling 22 OS=Homo sapiens GN=RGS22	7,00E-48
CU684871	1,00E-06	hypothetical protein DDB_G0280503 [Dictyostelium discoideum AX4]	9,00E-31
CU684892	1,00E-06	(sp Q9R0I6) Baculoviral IAP repeat-containing protein 4 OS=Rattus norvegicus GN=Xiap	5,00E-08
CU684896	1,00E-06	(sp Q5RA96) GMP synthase [glutamine-hydrolyzing] OS=Pongo abelii GN=GMPS	9,00E-93
CU684897	1,00E-06		
CU684901	1,00E-05	(sp Q9Y6R7) IgGFc-binding protein OS=Homo sapiens GN=FCGBP	3,00E-14
CU684921	1,00E-06		
CU684933	1,00E-06		
CU684937	1,00E-06		
CU684947	1,00E-06	(sp P28227) NADP-dependent malic enzyme OS=Anas platyrhynchos GN=ME1	5,00E-56
CU684955	1,00E-06	(sp P41570) 6-phosphogluconate dehydrogenase, decarboxylating OS=Ceratititis capitata GN=Pgd	8,00E-11
CU684967	1,00E-05		

CU684977	1,00E-06	(sp P25786) Proteasome subunit alpha type-1 OS=Homo sapiens GN=PSMA1	1,00E-114
CU684981	1,00E-05	(sp Q63HN8) RING finger protein 213 OS=Homo sapiens GN=RNF213	2,00E-16
CU684987	1,00E-06	(sp Q25147) Protein transport protein Sec61 subunit alpha OS=Halocynthia roretzi	9,00E-52
CU684993	1,00E-06	(sp A7Y2W8) Sodium- and chloride-dependent glycine transporter 1 OS=Xenopus laevis GN=slc6a9	1,00E-22
CU684998	1,00E-06		
CU685009	1,00E-06		
CU685019	1,00E-06	(sp Q91496) Proto-oncogene protein c-fos OS=Tetraodon fluviatilis GN=fos	5,00E-06
CU685023	1,00E-06		
CU685025	1,00E-06	(sp Q9H0K4) Radial spoke head-like protein 1 OS=Homo sapiens GN=RSHL1	1,00E-12
CU685044	1,00E-06		
CU685045	1,00E-06	(sp P52912) Nucleolysin TIA-1 OS=Mus musculus GN=Tia1	9,00E-20
CU685052	1,00E-05		
CU685057	1,00E-06	(sp Q6YHK3) CD109 antigen OS=Homo sapiens GN=CD109	3,00E-06
CU685062	1,00E-06	(sp Q92797) Symplekin OS=Homo sapiens GN=SYMPK	5,00E-49
CU685067	1,00E-06		
CU685080	1,00E-06	hypothetical protein BRAFLDRAFT_109858 [Branchiostoma floridae]	5,00E-06
CU685091	1,00E-06		
CU685093	1,00E-06	(sp P02553) Tubulin alpha chain (Fragment) OS=Lytechinus pictus	1,00E-41
CU685097	1,00E-06	(sp Q9VZF4) F-box/WD repeat-containing protein 7 OS=Drosophila melanogaster GN=ago	4,00E-13
CU685100	1,00E-06		
CU685110	1,00E-06	(sp Q9Y0Y2) Adenylosuccinate synthetase OS=Drosophila melanogaster GN=CG17273	2,00E-46
CU685116	1,00E-06		
CU685122	1,00E-06	hypothetical protein BRAFLDRAFT_124345 [Branchiostoma floridae]	2,00E-26
CU685125	1,00E-06	(sp Q8NHU2) Uncharacterized protein C20orf26 OS=Homo sapiens GN=C20orf26	1,00E-14
CU685130	1,00E-06	(sp P23098) Dynein beta chain, ciliary OS=Tripleneustes gratilla	2,00E-97
CU685134	1,00E-06		
CU685151	1,00E-05		

CU685153	1,00E-06	(sp Q5BJ29) F-box/LRR-repeat protein 7 OS=Mus musculus GN=Fbxl7	3,00E-16
CU685163	1,00E-06	(sp P05822) Transposase insF for insertion sequence IS3A/B/C/D/E/fA	2,00E-71
CU685194	1,00E-06	OS=Escherichia coli (strain K12)	
		(sp P54120) Protein AIG1 OS=Arabidopsis thaliana GN=AIG1	4,00E-07
CU685202	1,00E-06	(sp Q5T655) Coiled-coil domain-containing protein 147 OS=Homo sapiens	5,00E-49
CU685217	1,00E-06	GN=CCDC147	
CU685227	1,00E-06	(sp A5LFW4) EP300-interacting inhibitor of differentiation 3 OS=Macaca fascicularis GN=EID3	8,00E-26
CU685243	1,00E-06	(sp Q4R3Y4) Long-chain fatty acid transport protein 4 OS=Macaca fascicularis GN=SLC27A4	6,00E-59
CU685245	1,00E-06	(sp Q8BKT3) GC-rich sequence DNA-binding factor OS=Mus musculus	7,00E-06
CU685246	1,00E-06	GN=Tcf9	
CU685258	1,00E-06	(sp B2GV06) Succinyl-CoA:3-ketoacid-coenzyme A transferase 1, mitochondrial OS=Rattus norvegicus GN=Oxct1	2,00E-77
CU685260	1,00E-05		
CU685268	1,00E-05		
CU685269	1,00E-06	(sp P47791) Glutathione reductase, mitochondrial OS=Mus musculus GN=Gsr	8,00E-51
CU685270	1,00E-06		
CU685276	1,00E-06		
CU685281	1,00E-06		
CU685287	1,00E-06		
CU685296	1,00E-06	(sp P23469) Receptor-type tyrosine-protein phosphatase epsilon OS=Homo sapiens GN=PTPRE	1,00E-08
CU685297	1,00E-06	(sp P61255) 60S ribosomal protein L26 OS=Mus musculus GN=Rpl26	3,00E-50
CU685308	1,00E-06		
CU685330	1,00E-06	(sp Q9UHC1) DNA mismatch repair protein Mlh3 OS=Homo sapiens GN=MLH3	6,00E-11
CU685333	1,00E-06	IS10 transposase [Escherichia coli O157:H7 str. EC869]	1,00E-132
CU685341	1,00E-05	(sp P30337) N-chimaerin OS=Rattus norvegicus GN=Chn1	6,00E-20
CU685349	1,00E-06		
CU685368	1,00E-06		
CU685370	1,00E-06		

CU685377	1,00E-05		
CU685378	1,00E-06	(sp Q3MIS4) Uncharacterized protein C6orf224 OS=Homo sapiens	8,00E-29
CU685379	1,00E-06	GN=C6orf224	
CU685383	1,00E-06		
CU685398	1,00E-06		
CU685401	1,00E-06		
CU685426	1,00E-05	(sp P79024) Citrate synthase, mitochondrial OS=Candida tropicalis GN=CIT	3,00E-32
CU685431	1,00E-05		
CU685432	1,00E-06	(sp Q96F81) Protein dispatched homolog 1 OS=Homo sapiens GN=DISP1	2,00E-12
CU685436	1,00E-06	(sp O43390) Heterogeneous nuclear ribonucleoprotein R OS=Homo sapiens	7,00E-15
CU685437	1,00E-05	GN=HNRNPR	
CU685447	1,00E-06		
CU685451	1,00E-06	(sp A1L0Y2) Neuralized-like protein 4 OS=Xenopus tropicalis GN=neur14	9,00E-17
CU685453	1,00E-06		
CU685455	1,00E-06	(sp P57717) Estrogen receptor OS=Danio rerio GN=esr1	3,00E-75
CU685462	1,00E-06		
CU685480	1,00E-06		
CU685493	1,00E-05		
CU685496	1,00E-06		
CU685521	1,00E-06		
CU685526	1,00E-06		
CU685529	1,00E-06		
CU685532	1,00E-06		
CU685538	1,00E-06	(sp P98133) Fibrillin-1 OS=Bos taurus GN=FBN1	4,00E-26
CU685545	1,00E-06		
CU685546	1,00E-05		
CU685549	1,00E-06		
CU685550	1,00E-05		
CU685555	1,00E-06		
CU685559	1,00E-06		
CU685561	1,00E-06		
CU685577	1,00E-06	(sp Q27665) 6-phosphofructokinase OS=Haemonchus contortus GN=PFK	3,00E-74

CU685582	1,00E-06		
CU685599	1,00E-06		
CU685611	1,00E-06	(sp O54834) Rho GTPase-activating protein 6 OS=Mus musculus GN=Arhgap6	1,00E-08
CU685613	1,00E-06		
CU685636	1,00E-06		
CU685648	1,00E-06		
CU685652	1,00E-06		
CU685660	1,00E-06		
CU685661	1,00E-06	(sp Q6P4F7) Rho GTPase-activating protein 11A OS=Homo sapiens GN=ARHGAP11A	2,00E-16
CU685662	1,00E-06		
CU685666	1,00E-06	(sp P11182) Lipoamide acyltransferase component of branched-chain alpha- keto acid dehydrogenase complex	1,00E-21
CU685675	1,00E-06		
CU685677	1,00E-05	(sp P00548) Pyruvate kinase muscle isozyme OS=Gallus gallus GN=PKM2	5,00E-36
CU685686	1,00E-06	(sp Q60755) Calcitonin receptor OS=Mus musculus GN=Calcr	7,00E-11
CU685687	1,00E-06		
CU685690	1,00E-06	(sp Q32LE3) Centrin-1 OS=Bos taurus GN=CETN1	3,00E-52
CU685696	1,00E-06		
CU685698	1,00E-06		
CU685699	1,00E-06	(sp Q16270) Insulin-like growth factor-binding protein 7 OS=Homo sapiens GN=IGFBP7	6,00E-37
CU685701	1,00E-06		
CU685704	1,00E-06	(sp Q08420) Extracellular superoxide dismutase [Cu-Zn] OS=Rattus norvegicus GN=Sod3	6,00E-10
CU685708	1,00E-06		
CU685718	1,00E-06		
CU685719	1,00E-06		
CU685731	1,00E-06	(sp P82596) Perlucin OS=Haliotis laevigata	3,00E-08
CU685735	1,00E-06		
CU685749	1,00E-06		
CU685752	1,00E-06		
CU685765	1,00E-06	(sp Q9QW07) 1-phosphatidylinositol-4,5-bisphosphate phosphodiesterase beta-4 OS=Rattus norvegicus GN=Plcb4	3,00E-49

CU685769	1,00E-06	(sp P04113) Myosin regulatory light chain A, smooth adductor muscle OS=Patinopecten yessoensis	2,00E-61
CU685786	1,00E-06	(sp P11833) Tubulin beta chain OS=Paracentrotus lividus	1,00E-49
CU685796	1,00E-06	(sp P26642) Elongation factor 1-gamma-A OS=Xenopus laevis GN=eef1g-A	2,00E-40
CU685799	1,00E-05		
CU685802	1,00E-05	(sp O08605) MAP kinase-interacting serine/threonine-protein kinase 1 OS=Mus musculus GN=Mknk1	1,00E-112
CU685821	1,00E-06		
CU685844	1,00E-06	(sp P11507) Sarcoplasmic/endoplasmic reticulum calcium ATPase 2 OS=Rattus norvegicus GN=Atp2a2	8,00E-07
CU685866	1,00E-06	(sp P24733) Myosin heavy chain, striated muscle OS=Aequipecten irradians	1,00E-11
CU685894	1,00E-06	(sp Q26065) Actin, adductor muscle OS=Placopecten magellanicus (sp Q64319) Neutral and basic amino acid transport protein rBAT OS=Rattus norvegicus GN=Slc3a1	6,00E-32 1,00E-17
CU685917	1,00E-06		
CU685923	1,00E-06		
CU685927	1,00E-05		
CU685928	1,00E-06		
CU685937	1,00E-06		
CU685948	1,00E-06	(sp O96064) Paramyosin OS=Mytilus galloprovincialis	1,00E-29
CU685956	1,00E-05	(sp P50426) N-acetylglucosamine-6-sulfatase OS=Capra hircus GN=GNS	3,00E-08
CU685958	1,00E-06		
CU685965	1,00E-06	GF14191 [Drosophila ananassae]	4,00E-16
CU685974	1,00E-05		
CU685977	1,00E-06	(sp P30163) Actin-2 OS=Onchocerca volvulus GN=act-2b	0,00E+00
CU686034	1,00E-06		
CU686039	1,00E-06	(sp Q00232) NADH-ubiquinone oxidoreductase chain 5 OS=Mytilus edulis GN=ND5	7,00E-50
CU686044	1,00E-06	(sp Q9VU68) Actin-interacting protein 1 OS=Drosophila melanogaster GN=flr	2,00E-95
CU686063	1,00E-06	(sp O96064) Paramyosin OS=Mytilus galloprovincialis	3,00E-34
CU686086	1,00E-06		
CU686095	1,00E-06	(sp Q00232) NADH-ubiquinone oxidoreductase chain 5 OS=Mytilus edulis GN=ND5	7,00E-50

CU686096	1,00E-06	(sp P02637) Sarcoplasmic calcium-binding protein OS=Patinopecten yessoensis	7,00E-57
CU686111	1,00E-06	(sp Q78EG7) Protein tyrosine phosphatase type IVA 1 OS=Rattus norvegicus GN=Ptp4a1	3,00E-09
CU686124	1,00E-06	(sp A6NM10) Aquaporin-12B OS=Homo sapiens GN=AQP12B	7,00E-06
CU686142	1,00E-06	(sp P00399) Cytochrome c oxidase subunit 1 OS=Drosophila melanogaster GN=mt:Col	1,00E-137
CU686153	1,00E-06	(sp P26038) Moesin OS=Homo sapiens GN=MSN	2,00E-53
CU686185	1,00E-06		
CU686187	1,00E-06	(sp Q8CGF7) Transcription elongation regulator 1 OS=Mus musculus GN=Tcerg1	7,00E-47
CU686199	1,00E-06		
CU686227	1,00E-06	(sp P37805) Transgelin-3 OS=Rattus norvegicus GN=Tagln3	5,00E-08
CU686255	1,00E-06		
CU686259	1,00E-06		
CU686273	1,00E-06		
CU686275	1,00E-05	hypothetical protein [Schistosoma mansoni]	4,00E-64
CU686279	1,00E-06	(sp P24733) Myosin heavy chain, striated muscle OS=Aequipecten irradians	4,00E-25
CU686284	1,00E-06	(sp P48556) 26S proteasome non-ATPase regulatory subunit 8 OS=Homo sapiens GN=PSMD8	1,00E-84
CU686296	1,00E-06		
CU686300	1,00E-06	(sp P74897) Uncharacterized protein in QAH/OAS sulfhydrylase 3'region OS=Thermus aquaticus	1,00E-07
CU686302	1,00E-06		
CU686305	1,00E-06	(sp Q803E0) Paladin OS=Danio rerio GN=pald	3,00E-13
CU686339	1,00E-06	(sp P02886) Discoidin-1 subunit A OS=Dictyostelium discoideum GN=dscA-1	5,00E-06
CU686343	1,00E-06		
CU686381	1,00E-06		
CU686398	1,00E-05	(sp Q9VFP2) Protein roadkill OS=Drosophila melanogaster GN=rdx	1,00E-163
CU686403	1,00E-06	(sp P10184) Ovoinhibitor OS=Gallus gallus GN=OIH	5,00E-27
CU686433	1,00E-05		
CU686462	1,00E-06	(sp Q8VCD7) Lysine-specific demethylase 4C OS=Mus musculus GN=Kdm4c	6,00E-67

CU686467	1,00E-06	(sp P24051) 40S ribosomal protein S27-like protein OS=Rattus norvegicus	
CU686468	1,00E-06	GN=Rps27l	2,00E-40
CU686478	1,00E-06	(sp P37397) Calponin-3 OS=Rattus norvegicus GN=Cnn3	7,00E-31
CU686480	1,00E-06	Trypanosoma brucei TREU927 transposase of Tn10 partial mRNA	0,00E+00
CU686482	1,00E-06	(sp P90689) Actin OS=Brugia malayi	1,00E-32
CU686483	1,00E-06	(sp B5FZ63) Enhancer of yellow 2 transcription factor homolog	
CU686485	1,00E-06	OS=Taeniopygia guttata GN=ENY2	1,00E-34
CU686511	1,00E-06	(sp Q4PMB3) 40S ribosomal protein S4 OS=Ixodes scapularis GN=RpS4	1,00E-119
CU686533	1,00E-05	(sp P62919) 60S ribosomal protein L8 OS=Rattus norvegicus GN=Rpl8	1,00E-124
CU686553	1,00E-06		
CU686554	1,00E-06	(sp Q7ZUY3) Histone H2A.x OS=Danio rerio GN=h2afx	2,00E-38
CU686568	1,00E-06		
CU686569	1,00E-06	(sp P10184) Ovoinhibitor OS=Gallus gallus GN=OIH	5,00E-27
CU686580	1,00E-06	(sp P49395) 40S ribosomal protein S3a OS=Aplysia californica GN=RPS3A	1,00E-107
CX068896	1,00E-06	(sp P00399) Cytochrome c oxidase subunit 1 OS=Drosophila melanogaster	
CX068897	1,00E-06	GN=mt:Col	1,00E-137
CX068898	1,00E-06	(sp P00399) Cytochrome c oxidase subunit 1 OS=Drosophila melanogaster	
CX068901	1,00E-06	GN=mt:Col	1,00E-137
CX068907	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00
CX068909	1,00E-06	(sp Q24940) Cathepsin L-like proteinase OS=Fasciola hepatica GN=Cat-1	4,00E-15
CX068911	1,00E-06	(sp P24051) 40S ribosomal protein S27-like protein OS=Rattus norvegicus	
CX068914	1,00E-06	GN=Rps27l	2,00E-40
CX068918	1,00E-06		
CX068919	1,00E-06	(sp Q3T0Q6) Cellular nucleic acid-binding protein OS=Bos taurus GN=CNBP	2,00E-20
CX068920	1,00E-06	(sp Q9YIC0) Elongation factor 1-alpha OS=Oryzias latipes GN=eef1a	0,00E+00
CX068923	1,00E-06	Drosophila willistoni GK25336 (DwilGK25336), mRNA	3,00E-06
CX068925	1,00E-06	(sp P00399) Cytochrome c oxidase subunit 1 OS=Drosophila melanogaster	
		GN=mt:Col	1,00E-137

CX068933	1,00E-06		
CX068934	1,00E-06	(sp Q9ZRB7) Tubulin alpha chain OS=Triticum aestivum GN=TUBA	2,00E-06
CX068935	1,00E-05		
CX068939	1,00E-06		
CX068943	1,00E-06		
CX068946	1,00E-05	(sp Q02105) Complement C1q subcomponent subunit C OS=Mus musculus GN=C1qc	3,00E-06
CX068947	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00
CX068955	1,00E-06	(sp P30044) Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5 (sp P00399) Cytochrome c oxidase subunit 1 OS=Drosophila melanogaster GN=mt:Col	5,00E-52 1,00E-137
CX068961	1,00E-06		
CX068964	1,00E-06		
CX068975	1,00E-06		
CX068978	1,00E-06		
CX068988	1,00E-06		
CX068992	1,00E-06		
CX068993	1,00E-06		
CX068996	1,00E-06		
CX068997	1,00E-06	PREDICTED: hypothetical protein isoform 1 [Ciona intestinalis]	8,00E-12
CX068999	1,00E-05		
CX069000	1,00E-06		
CX069002	1,00E-06	hypothetical protein IscW_ISCW009032 [Ixodes scapularis]	3,00E-07
CX069006	1,00E-06		
CX069008	1,00E-06		
CX069016	1,00E-06	hypothetical protein BRAFLDRAFT_71169 [Branchiostoma floridae]	7,00E-09
CX069020	1,00E-06	(sp P30163) Actin-2 OS=Onchocerca volvulus GN=act-2b	0,00E+00
CX069022	1,00E-06		
CX069026	1,00E-06	(sp P00399) Cytochrome c oxidase subunit 1 OS=Drosophila melanogaster GN=mt:Col	1,00E-137
CX069027	1,00E-05	(sp Q26065) Actin, adductor muscle OS=Placopecten magellanicus	0,00E+00
CX069030	1,00E-06		
CX069031	1,00E-06		
CX069033	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00
CX069034	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00

CX069035	1,00E-06	(sp P30044) Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5	5,00E-52
CX069038	1,00E-05		
CX069040	1,00E-06	(sp Q6Y263) 60S ribosomal protein L24 OS=Pagrus major GN=rpl24	9,00E-41
CX069041	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00
CX069047	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00
CX069052	1,00E-06		
CX069053	1,00E-06		
CX069058	1,00E-06		
CX069059	1,00E-06		
CX069063	1,00E-06		
CX069065	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00
CX069067	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00
CX069069	1,00E-06		
CX069073	1,00E-06		
CX069076	1,00E-05		
CX069077	1,00E-06		
CX069078	1,00E-06		
CX069082	1,00E-06		
CX069083	1,00E-06		
CX069084	1,00E-06	(sp P07291) Myosin essential light chain, striated adductor muscle OS=Aequipecten irradians	1,00E-56
CX069090	1,00E-06		
CX069091	1,00E-06		
CX069092	1,00E-06		
CX069095	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00
CX069097	1,00E-06	(sp Q9XP77) Cytochrome b OS=Sminthopsis ooldea GN=MT-CYB	5,00E-70
CX069098	1,00E-06		
CX069104	1,00E-05	(sp P06603) Tubulin alpha-1 chain OS=Drosophila melanogaster GN=alphaTub84B	1,00E-173
CX069105	1,00E-06		
CX069106	1,00E-06	(sp Q9BXS5) AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1	0,00E+00
CX069107	1,00E-06	(sp Q9BYV8) Centrosomal protein of 41 kDa OS=Homo sapiens GN=TSGA14	2,00E-17

CX069111	1,00E-06	(sp Q9QYP1) Low-density lipoprotein receptor-related protein 4 OS=Rattus norvegicus GN=Lrp4	2,00E-30
CX069216	1,00E-06	(sp P62329) Thymosin beta-4 OS=Rattus norvegicus GN=Tmsb4x	1,00E-12
CX069218	1,00E-06	(sp O61463) 60S acidic ribosomal protein P2 OS=Cryptochiton stelleri	1,00E-17
CX069220	1,00E-06	(sp Q5RC38) Cytochrome c oxidase subunit 6A2, mitochondrial OS=Pongo abelii GN=COX6A2	2,00E-06
CX069221	1,00E-06		
CX069225	1,00E-06	(sp O61463) 60S acidic ribosomal protein P2 OS=Cryptochiton stelleri	1,00E-17
CX069226	1,00E-06	(sp Q02543) 60S ribosomal protein L18a OS=Homo sapiens GN=RPL18A	3,00E-69
CX069230	1,00E-06	(sp Q7PCJ9) Diamine acetyltransferase 2 OS=Sus scrofa GN=SAT2	8,00E-19
CX069232	1,00E-05	(sp Q6DDU9) RNA-binding protein 5-B OS=Xenopus laevis GN=rmb5-B	2,00E-23
CX069245	1,00E-06	(sp P06603) Tubulin alpha-1 chain OS=Drosophila melanogaster GN=alphaTub84B	0,00E+00
CX069254	1,00E-06	(sp Q6YHK3) CD109 antigen OS=Homo sapiens GN=CD109	8,00E-55
CX069255	1,00E-05	(sp Q86VB7) Scavenger receptor cysteine-rich type 1 protein M130 OS=Homo sapiens GN=CD163	1,00E-26
CX069260	1,00E-06		
CX069261	1,00E-06		
CX069262	1,00E-06	(sp Q4PMB3) 40S ribosomal protein S4 OS=Ixodes scapularis GN=RpS4	1,00E-119
CX069268	1,00E-06	(sp Q6P878) Eukaryotic translation initiation factor 3 subunit L OS=Xenopus tropicalis GN=eif3l	1,00E-100
CX069271	1,00E-06	(sp Q2M389) UPF0681 protein KIAA1033 OS=Homo sapiens GN=KIAA1033	7,00E-56
CX069277	1,00E-06	(sp P92958) SNF1-related protein kinase catalytic subunit alpha KIN11 OS=Arabidopsis thaliana GN=KIN11	7,00E-09
CX069279	1,00E-06	(sp P04069) Carboxypeptidase B OS=Astacus fluviatilis	2,00E-57
CX069282	1,00E-06	(sp Q26636) Cathepsin L OS=Sarcophaga peregrina	1,00E-102
CX069284	1,00E-06	hypothetical protein sce4070 [Sorangium cellulosum 'So ce 56']	2,00E-45
CX069285	1,00E-06	(sp Q9QYP1) Low-density lipoprotein receptor-related protein 4 OS=Rattus norvegicus GN=Lrp4	2,00E-30
CX069286	1,00E-06	(sp P62972) Ubiquitin OS=Xenopus laevis	5,00E-36
CX069287	1,00E-06	(sp Q26065) Actin, adductor muscle OS=Placopecten magellanicus	0,00E+00
CX069289	1,00E-06	(sp P15253) Calreticulin OS=Oryctolagus cuniculus GN=CALR	1,00E-124
CX069290	1,00E-06	(sp P68372) Tubulin beta-2C chain OS=Mus musculus GN=Tubb2c	0,00E+00
CX069293	1,00E-06	(sp O00339) Matrilin-2 OS=Homo sapiens GN=MATN2	1,00E-14

CX069294	1,00E-06	(sp Q01518) Adenylyl cyclase-associated protein 1 OS=Homo sapiens GN=CAP1 SV=4	2,00E-47
CX069297	1,00E-06	(sp P28840) Neuroendocrine convertase 1 OS=Rattus norvegicus GN=Pcsk1	9,00E-52
CX069303	1,00E-05	(sp Q922J9) Fatty acyl-CoA reductase 1 OS=Mus musculus GN=Far1	1,00E-142
CX069306	1,00E-06	(sp P12261) Elongation factor 1-gamma OS=Artemia salina	1,00E-112
CX069310	1,00E-06	(sp A2AX52) Collagen alpha-4(VI) chain OS=Mus musculus GN=Col6a4	2,00E-13
CX069319	1,00E-06	hypothetical protein TRIADDRAFT_7530 [Trichoplax adhaerens] (sp POC6B8) Sushi, von Willebrand factor type A, EGF and pentraxin domain- containing protein 1	7,00E-06
CX069321	1,00E-06	(sp P08297) Early nodulin-75 OS=Glycine max GN=ENOD2A	4,00E-07
CX069323	1,00E-06	(sp Q6P823) Histone H3.3 OS=Xenopus tropicalis GN=TGas113e22.1	3,00E-14
CX069324	1,00E-06		5,00E-70
CX069326	1,00E-06	(sp A7EGL7) ATP-dependent RNA helicase eIF4A OS=Sclerotinia sclerotiorum	3,00E-54
CX069331	1,00E-06	(sp Q6PC69) 60S ribosomal protein L10a OS=Danio rerio GN=rpl10a	1,00E-67
CX069333	1,00E-06	(sp P02434) 60S ribosomal protein L35a OS=Xenopus laevis GN=rpl35a	9,00E-41
CX069337	1,00E-06		
CX069340	1,00E-06	(sp P25867) Ubiquitin-conjugating enzyme E2-17 kDa OS=Drosophila melanogaster GN=eff	8,00E-64
CX069347	1,00E-06	(sp P41824) Y-box factor homolog OS=Aplysia californica	2,00E-39
CX069348	1,00E-06	(sp Q5M8V0) Transcription factor BTF3 homolog 4 OS=Xenopus tropicalis GN=btf3l4	2,00E-52
FP089714	1,00E-06	(sp P40618) High mobility group protein B3 OS=Gallus gallus GN=HMGB3	1,00E-11
FP089731	1,00E-06		
FP089746	1,00E-06		
FP089757	1,00E-06	(sp Q98893) Vitellogenin-2 OS=Fundulus heteroclitus	1,00E-10
FP089760	1,00E-06	hypothetical protein BRAFLDRAFT_125849 [Branchiostoma floridae] (sp Q5T2S8) Armadillo repeat-containing protein 4 OS=Homo sapiens GN=ARMC4	2,00E-09
FP089761	1,00E-06		3,00E-97
FP089763	1,00E-06	(sp P52742) Zinc finger protein 135 OS=Homo sapiens GN=ZNF135	3,00E-38
FP089766	1,00E-06	ATP synthase FO subunit 6 [Crassostrea gigas]	1,00E-105
FP089773	1,00E-06	(sp Q7A185) Protein fmtA OS=Staphylococcus aureus (strain MW2) GN=fmtA	8,00E-06
FP089776	1,00E-06	(sp Q98893) Vitellogenin-2 OS=Fundulus heteroclitus	7,00E-07

FP089783	1,00E-06	(sp Q8TGM7) Uncharacterized protein ART2 OS= <i>Saccharomyces cerevisiae</i> GN=ART2 PE=4	2,00E-16
FP089803	1,00E-06	(sp Q69ZN7) Myoferlin OS= <i>Mus musculus</i> GN=Myof	1,00E-142
FP089813	1,00E-06	(sp P06857) Pancreatic lipase-related protein 1 OS= <i>Canis familiaris</i> GN=PNLIPRP1	1,00E-18
FP089816	1,00E-06	(sp Q61941) NAD(P) transhydrogenase, mitochondrial OS= <i>Mus musculus</i> GN=Nnt	2,00E-54
FP089817	1,00E-06		
FP089856	1,00E-06	<i>Toxoplasma gondii</i> ME49 hypothetical protein, mRNA	1,00E-06
FP089872	1,00E-06	(sp Q9W638) Zinc finger protein ubi-d4 A OS= <i>Xenopus laevis</i> GN=req-A	3,00E-43
FP089900	1,00E-05		
FP089901	1,00E-06		
FP089904	1,00E-06	(sp P60755) MAM domain-containing glycosylphosphatidylinositol anchor protein 2 OS= <i>Mus musculus</i> GN=Mdga2	5,00E-18
FP089916	1,00E-06	(sp Q4R7Y5) DnaJ homolog subfamily B member 6 OS= <i>Macaca fascicularis</i> GN=DNAJB6	1,00E-10
FP089929	1,00E-06		
FP089937	1,00E-06		
FP089949	1,00E-06	hypothetical protein BRAFLDRAFT_117068 [<i>Branchiostoma floridae</i>]	7,00E-07
FP090914	1,00E-06		
FP090926	1,00E-06	(sp Q62288) Testican-1 OS= <i>Mus musculus</i> GN=Spock1	6,00E-30
FP090929	1,00E-06	(sp A7Y2X0) Sodium- and chloride-dependent glycine transporter 2 OS= <i>Xenopus laevis</i> GN=slc6a5	1,00E-40
FP090965	1,00E-06		
FP090972	1,00E-06		
FP090992	1,00E-06	(sp Q17N71) Protein KIAA0664 homolog OS= <i>Aedes aegypti</i> GN=AAEL000794	8,00E-48
FP090996	1,00E-06	predicted protein [<i>Laccaria bicolor</i> S238N-H82]	1,00E-29
FP090998	1,00E-06	(sp Q8TGM7) Uncharacterized protein ART2 OS= <i>Saccharomyces cerevisiae</i> GN=ART2 PE=4	2,00E-16
FP091003	1,00E-06		
FP091015	1,00E-06		
FP091019	1,00E-06		
FP091029	1,00E-06	hypothetical protein BRAFLDRAFT_87842 [<i>Branchiostoma floridae</i>]	6,00E-08

FP091045	1,00E-06	(sp Q64512) Tyrosine-protein phosphatase non-receptor type 13 OS=Mus musculus GN=Ptpn13	4,00E-29
FP091048	1,00E-06	(sp A2VDZ3) Myocyte-specific enhancer factor 2A OS=Bos taurus GN=MEF2A	3,00E-46
FP091051	1,00E-06		
FP091055	1,00E-06	(sp A7EGL7) ATP-dependent RNA helicase eIF4A OS=Sclerotinia sclerotiorum	3,00E-54
FP091067	1,00E-06		
FP091118	1,00E-06		
FP091130	1,00E-06	(sp P46088) Glutathione S-transferase OS=Ommastrephes sloanei	2,00E-32
FP091155	1,00E-06		
FP091157	1,00E-05	(sp Q6RWA9) Sodium/potassium-transporting ATPase subunit alpha OS=Taenia solium	3,00E-49
FP091177	1,00E-06		
FP091188	1,00E-06		
FP091193	1,00E-06		
FP091213	1,00E-06		
Y08370	1,00E-06		
