

Genbank	TC	Log2(GM ratio)	Role_guess
C88186	TC411856	-3.391007323	weakly similar to cytosolic phospholipase A2 gamma - human, partial (47%)
C86518		-3.320587673	
AU045621	TC389983	-3.156487079	
C78577	TC429425	-1.953107529	similar to lactase-phlorizin hydrolase precursor (Rattus rattus), partial (9%)
C85356	TC379554	-1.849192326	
AW537044	TC375072	-1.65248673	Unknown (protein for MGC:6302) (Mus musculus), complete
C79107	TC382580	-1.613010585	putative (Mus musculus), complete
AU043044	TC401153	-1.60039748	ARGINASE II PRECURSOR (EC 3.5.3.1) (NON-HEPATIC ARGINASE) (KIDNEY-TYPE ARGINASE). [Mouse], complete
AU017156		-1.560720476	
A1850731	TC382580	-1.539770974	putative (Mus musculus), complete
AW546196	TC425599	-1.506761575	neutral ceramidase (Mus musculus), complete
AU040351	TC375368	-1.490288994	CARBONIC ANHYDRASE IV PRECURSOR (EC 4.2.1.1) (CARBONATE DEHYDRATASE IV) (CA-IV). [Mouse], complete
A1838895	TC404479	-1.434664926	AROMATIC-L-AMINO-ACID DECARBOXYLASE (EC 4.1.1.28) (AADC) (DOPA DECARBOXYLASE) (DDC). [Mouse], partial (27%)
A1843134	TC404488	-1.41873821	homologue to HYPOTHETICAL PROTEIN KIAA0205. [Human], complete
C66592	TC374820	-1.406974822	homologue to aldolase B (Rattus norvegicus), complete
A1840448	TC375055	-1.40654839	homologue to hypothetical protein (Macaca fascicularis), partial (52%)
AW537122	TC425440	-1.406405474	homologue to ba11D8.2.1 (novel protein (possible GTP binding protein) (isoform 1)) (Homo sapiens), complete
AU043077	TC378903	-1.371788845	family 4 cytochrome P450 (Mus musculus), complete
BE380947	TC389966	-1.361934922	ANGIOTENSIN-CONVERTING ENZYME, TESTIS-SPECIFIC ISOFORM PRECURSOR (EC 3.4.15.1) (ACE-T) (DIPEPTIDYL CARBOXYPEPTIDASE I) (KININASE II). [Mouse], complete
C81194	TC425659	-1.34369524	heat shock protein 105 kDa alpha (Mus musculus), complete
AU016367	TC404488	-1.333849944	homologue to HYPOTHETICAL PROTEIN KIAA0205. [Human], complete
A1835590	TC425440	-1.333401212	homologue to ba11D8.2.1 (novel protein (possible GTP binding protein) (isoform 1)) (Homo sapiens), complete
AU045090	TC389362	-1.324254255	
A1853000	TC443762	-1.301867895	Similar to glutathione S-transferase, theta 1 (Mus musculus), complete
A1845853	TC425202	-1.278724192	Similar to fibulin 1 (Mus musculus), complete
AW545641	TC394270	-1.274654255	homologue to putative (Mus musculus), partial (53%)
AA400683	TC397097	-1.265968256	
AW544862	TC328235	-1.241262954	homologue to heat shock protein 105 kDa beta (42 degrees C-specific heat shock protein) (Mus musculus), complete
A1852543	TC397002	-1.235732406	SOLUBLE EPOXIDE HYDROLASE (SEH) (EC 3.3.2.3) (EPOXIDE HYDRATASE) (CYTOSOLIC EPOXIDE HYDROLASE) (CEH). [Mouse], complete
C87836	TC368558	-1.225573288	putative (Mus musculus), complete
AW538067	TC400374	-1.213976284	CREATINE KINASE, B CHAIN (EC 2.7.3.2) (B-CK). [Mouse], complete
AW536153	TC393994	-1.208121912	MICROSOMAL TRIGLYCERIDE TRANSFER PROTEIN, LARGE SUBUNIT PRECURSOR. [Mouse], complete
AW545252	TC386181	-1.206784491	RIKEN cDNA 1110029G07 gene (Mus musculus), complete
A1838813		-1.198373711	
A1845580	TC386181	-1.194234122	RIKEN cDNA 1110029G07 gene (Mus musculus), complete
A1838867	TC420975	-1.180330557	malate dehydrogenase (EC 1.1.1.37), cytosolic - mouse, complete
AU015411	TC421977	-1.172797084	GUANINE DEAMINASE (EC 3.5.4.3) (GUANASE) (GUANINE AMINOHYDROLASE) (GAH). [Mouse], complete
AU022217	TC397002	-1.172623848	SOLUBLE EPOXIDE HYDROLASE (SEH) (EC 3.3.2.3) (EPOXIDE HYDRATASE) (CYTOSOLIC EPOXIDE HYDROLASE) (CEH). [Mouse], complete
AU022388	TC393392	-1.171159331	homologue to tripartite motif protein TRIM30 isoform alpha (Mus musculus), partial (48%)
AW536474	TC435596	-1.161361049	apolipoprotein A-IV (Mus musculus castaneus), complete
A1838228	TC440242	-1.152388244	leukemia/lymphoma related factor LRF (Mus musculus), complete
AW557879	TC421186	-1.141446533	PDZK1 (Mus musculus), complete
C78825	TC421186	-1.140086133	PDZK1 (Mus musculus), complete
AU040965	TC432346	-1.13483355	similar to aldose 1-epimerase (Sus scrofa), complete
A1849532	TC397418	-1.133940709	putative (Mus musculus), complete
AU016472	TC379400	-1.122795631	similar to unknown (Homo sapiens), partial (97%)
BE373153	TC435490	-1.106284514	stearyl-Coenzyme A desaturase 1 (Mus musculus), complete
AW543802	TC425686	-1.103727307	DEOXYRIBONUCLEASE GAMMA PRECURSOR (EC 3.1.21.-) (DNASE GAMMA) (DEOXYRIBONUCLEASE I-LIKE 3) (DNASE I HOMOLOGOUS PROTEIN DHP2) (LIVER AND SPLEEN DNASE) (LS-DNASE) (LSD). [Mouse], complete
A1851917	TC379400	-1.101472393	similar to unknown (Homo sapiens), partial (97%)
C86683	TC414908	-1.094701245	
C80512	TC417612	-1.087601094	CREATINE KINASE, UBIQUITOUS MITOCHONDRIAL PRECURSOR (EC 2.7.3.2) (U-MTCK) (MIA-CK) (ACIDIC-TYPE MITOCHONDRIAL CREATINE KINASE). [Mouse], complete
AW557809	TC368153	-1.08050764	similar to KIAA1564 protein (Homo sapiens), partial (84%)
A1834402	TC393642	-1.08059112	MAD PROTEIN (MAX DIMERIZER). [Mouse], complete
A1852832	TC379668	-1.066230096	T-CELL ANTIGEN CD7 PRECURSOR. [Mouse], complete
AU016275	TC368434	-1.061622367	APOA1N PRECURSOR (EC 3.4.22.-) (CYSTEINE PROTEASE CPP32) (YAMA PROTEIN) (CPP-32) (CASPASE-3) (CASP-3) (SREBP CLEAVAGE ACTIVITY 1) (SCA-1) (LICE). [Mouse], complete
AU040822	TC378830	-1.060021635	putative (Mus musculus), complete
C87210	TC371459	-1.056257227	putative (Mus musculus), complete
AU041673	TC439386	-1.050754337	malic enzyme, supematant (Mus musculus), complete
A1850687	TC368101	-1.047989991	putative (Mus musculus), complete
AW413200	TC400374	-1.046727313	CREATINE KINASE, B CHAIN (EC 2.7.3.2) (B-CK). [Mouse], complete
A1853466	TC401149	-1.045355167	putative (Mus musculus), complete
A1849037	TC393780	-1.045168638	LARGE NEUTRAL AMINO ACIDS TRANSPORTER SMALL SUBUNIT 2 (L-TYPE AMINO ACID TRANSPORTER 2). [Mouse], complete
A1836830	TC371459	-1.042731056	putative (Mus musculus), complete
AU022597	TC390619	-1.034145349	
A1843665	TC404393	-1.032394833	PROTEIN TRANSPORT PROTEIN SEC23A (SEC23-RELATED PROTEIN A). [Human], complete
AU016778	TC390056	-1.029367566	similar to carboxylesterase RL1 (Rattus norvegicus), complete
A1854179	TC432154	-1.028755752	similar to Similar to CGI-45 protein (Homo sapiens), partial (86%)
C79756	TC436439	-1.018665573	ALKALINE PHOSPHATASE, EMBRYONIC PRECURSOR (EC 3.1.3.1) (EAP). [Mouse], complete
AW558234	TC382592	-1.017918189	SESTRIN 1 (P53-REGULATED PROTEIN PA26). [Mouse], complete
AW558053	TC382201	-1.015769205	UDP-GLUCURONOSYLTRANSFERASE 1-2 PRECURSOR, MICROSOMAL (EC 2.4.1.17) (UDPGT) (UGT1*0) (UGT1-02) (UGT1.2) (UGT1A2) (BILIRUBIN SPECIFIC). [Mouse], complete
A1835775	TC378520	-1.014106087	homologue to ATP specific succinyl CoA synthetase beta subunit precursor (Homo sapiens), complete
A1842877	TC422120	-1.011026003	similar to glutathione S-transferase mu-class subunit M1 (Macaca fascicularis), complete
C78412	TC425440	-1.009803594	homologue to ba11D8.2.1 (novel protein (possible GTP binding protein) (isoform 1)) (Homo sapiens), complete
AU021475	TC432520	-1.00767357	homologue to myosin light chain kinase (Homo sapiens), partial (29%)
AU020215	TC424654	-1.005643811	SORBITOL DEHYDROGENASE (EC 1.1.1.14) (L-IDITOL 2-DEHYDROGENASE) (FRAGMENT). [Mouse], complete
AU018948	TC397356	-1.005080586	PROLACTIN RECEPTOR PRECURSOR (PRL-R). [Mouse], complete
A1841400	TC407907	-1.001264711	similar to unnamed protein product (Homo sapiens), complete
A1838930	TC367832	-0.999812555	Unknown (protein for MGC:6272) (Mus musculus), complete
A1842442		-0.995625846	
C85570	TC420975	-0.993564875	malate dehydrogenase (EC 1.1.1.37), cytosolic - mouse, complete
A1838757	TC428803	-0.986146023	homologue to AMINE OXIDASE [FLAVIN-CONTAINING] A (EC 1.4.3.4) (MONOAMINE OXIDASE) (MAO-A). [Rat], complete
AW557135	TC375114	-0.97806079	xanthine dehydrogenase (Mus musculus), complete
AW544861	TC409479	-0.97641399	
A1839122	TC368136	-0.974482088	DIMETHYLANILINE MONOOXYGENASE [N-OXIDE FORMING] 1 (EC 1.14.13.8) (HEPATIC FLAVIN-CONTAINING MONOOXYGENASE 1) (FMO 1) (DIMETHYLANILINE OXIDASE 1). [Mouse], complete
C77059	TC425098	-0.97234749	putative (Mus musculus), complete
A1848955	TC443482	-0.968618681	similar to unnamed protein product (Homo sapiens), complete
A1835380	TC367651	-0.967289035	putative (Mus musculus), complete
C87907	TC420975	-0.960697137	malate dehydrogenase (EC 1.1.1.37), cytosolic - mouse, complete
AU040826	TC421233	-0.957757603	homologue to LONG-CHAIN-FATTY-ACID-COA LIGASE 5 (EC 6.2.1.3) (LONG-CHAIN ACYL-COA SYNTHETASE 5) (LACS 5). [Rat], complete
A1838612	TC397129	-0.955470169	glutathione transferase (EC 2.5.1.18) mu2 - mouse, complete
A1850729		-0.952489592	

AI843123		-0.952192292
C86052	TC386307	-0.949886307 CALPONIN H1, SMOOTH MUSCLE (BASIC CALPONIN) (CALPONIN 1). [Mouse], complete
AI850717	TC385896	-0.949461809 putative (Mus musculus), complete
AW544700	TC417925	-0.947684943 FYVE zinc finger phosphatase (Mus musculus), complete
C80039	TC414006	-0.945150928 aspartate aminotransferase (Mus musculus), complete
AI849517	TC409010	-0.943165423 homologue to CHOLINESTERASE PRECURSOR (EC 3.1.1.8) (ACYLCHOLINE ACYLHYDROLASE) (CHOLINE ESTERASE II) (BUTYRYLCHOLINE ESTERASE) (PSEUDOCHELINESTERASE). [Mouse], partial (27%)
AI845946	TC382435	-0.942863276 homologue to heat shock protein 105 kDa beta (42 degrees C-specific heat shock protein) (Mus musculus), complete
AI850689	TC390450	-0.942381911 weakly similar to KIAA0443 (Homo sapiens), partial (42%)
AI853207	TC421305	-0.941666371
C80388	TC421860	-0.940502489 PROTEIN KINASE C, ZETA TYPE [EC 2.7.1.-] (NPKC-ZETA). [Mouse], complete
AW555156	TC422120	-0.938189902 similar to glutathione S-transferase mu-class subunit M1 (Macaca fascicularis), complete
AU021897	TC410869	-0.937261386
AW552780	TC413767	-0.93275325 glutathione transferase (EC 2.5.1.18) mu1 - mouse, complete
AA409867	TC382139	-0.930538512 cytochrome c [validated] - mouse, complete
AI842900	TC436506	-0.92954952
AW558952	TC443954	-0.928303414 transcription factor GATA-4 (Mus musculus), complete
AI854647	TC421923	-0.927696997 cationic amino acid transporter (Mus musculus), complete
AU046150	TC424846	-0.927566586 homologue to putative (Mus musculus), complete
AI843118	TC383098	-0.923996615 GAMMA-AMINOBUTYRIC ACID TYPE B RECEPTOR, SUBUNIT 1 PRECURSOR (GABA-B RECEPTOR 1) (GABA-B-R1) (GB1). [Mouse], complete
AW553206	TC413767	-0.923104554 glutathione transferase (EC 2.5.1.18) mu1 - mouse, complete
AW537347	TC428803	-0.92308695 homologue to AMINE OXIDASE [FLAVIN-CONTAINING] A (EC 1.4.3.4) (MONOAMINE OXIDASE) (MAO-A). [Rat], complete
AI838755	TC397890	-0.921615265
AI838594	TC414427	-0.918563908 homologue to Similar to PTD015 protein (Homo sapiens), complete
AW539541	TC371459	-0.918183505 putative (Mus musculus), complete
AI843740	TC417925	-0.917987849 FYVE zinc finger phosphatase (Mus musculus), complete
AU022525	TC393780	-0.915698734 LARGE NEUTRAL AMINO ACIDS TRANSPORTER SMALL SUBUNIT 2 (L-TYPE AMINO ACID TRANSPORTER 2). [Mouse], complete
AI853409	TC396816	-0.907775632 homologue to tricarboxylate carrier - rat (fragment), complete
AU021466	TC424654	-0.90769751 SORBITOL DEHYDROGENASE (EC 1.1.1.14) (L-IDITOL 2-DEHYDROGENASE) (FRAGMENT). [Mouse], complete
AI853530	TC421586	-0.904420232 similar to Unknown (protein for MGC:10733) (Homo sapiens), partial (58%)
AW544751	TC385920	-0.904296902 nucleoside triphosphatase (Mus musculus), complete
AW542441	TC390278	-0.904234009 Similar to acid sphingomyelinase-like phosphodiesterase (Mus musculus), complete
BE377125	TC393936	-0.900121203 malic enzyme, supernatant (Mus musculus), complete
AA407103	TC443288	-0.896240406 homologue to Unknown (protein for IMAGE:3509929) (Homo sapiens), partial (95%)
AW552108	TC382143	-0.887630034 putative (Mus musculus), complete
AI836560	TC420992	-0.887560664 transcobalamin II (Mus musculus), complete
AI852173	TC431920	-0.883248201 ORNITHINE DECARBOXYLASE ANTIZYME INHIBITOR. [Mouse], complete
AI839897	TC422087	-0.88292985
AI852898	TC372111	-0.879262949 putative (Mus musculus), complete
AI849776	TC386411	-0.877499798 Similar to RIKEN cDNA 2210023K21 gene (Mus musculus), complete
AI842323	TC436061	-0.872934656 neuronal IL-16 (Mus musculus), complete
AW536138	TC396566	-0.870778302 glucosamine-6-phosphate deaminase (Mus musculus), complete
AA410071	TC405707	-0.867889028
AI853879	TC436407	-0.86697172 SMAR1 (Mus musculus), complete
AI848096	TC414824	-0.865020422 protein 4.1B (Mus musculus), complete
C87789	TC401910	-0.864269999
AI849373	TC382139	-0.859822811 cytochrome c [validated] - mouse, complete
AI852055	TC425515	-0.8575229 mDab555 protein (Mus musculus), partial (55%)
AI840451	TC413776	-0.857260469 4F2 CELL SURFACE ANTIGEN HEAVY CHAIN (4F2HC). [Mouse], complete
AW536815	TC429067	-0.852235213 URIDINE PHOSPHORYLASE (EC 2.4.2.3) (UDRPASE). [Mouse], complete
AI846650	TC404668	-0.851152365 serotonin transporter (Mus musculus), complete
C85601		-0.85029447
AI844801	TC421010	-0.849556638 Unknown (protein for IMAGE:3981915) (Mus musculus), complete
AU021079	TC421950	-0.84831693 homologue to transmembrane protein induced by tumor necrosis factor alpha (Homo sapiens), complete
AI847132		-0.845892572
AW538333	TC374629	-0.844077295 homologue to SODIUM/POTASSIUM-TRANSPORTING ATPASE ALPHA-1 CHAIN PRECURSOR (EC 3.6.3.9) (SODIUM PUMP) (NA+K+ ATPASE). [Rat], complete
AI848937	TC400310	-0.841720277 putative (Mus musculus), complete
AW547940	TC368015	-0.841272334 similar to hypothetical protein KIAA0232 (Homo sapiens), partial (21%)
AI845780	TC404917	-0.838959052 tetratricopeptide repeat-containing protein - mouse, complete
AI843055	TC443605	-0.837948925 similar to ETS-RELATED TRANSCRIPTION FACTOR ELF-1 (E74-LIKE FACTOR 1). [Mouse], partial (67%)
AU022893	TC397001	-0.836339443 UDP-GLUCOSE 6-DEHYDROGENASE (EC 1.1.1.22) (UDP-GLC DEHYDROGENASE) (UDP-GLCDH) (UDPGDH). [Mouse], complete
AI838739	TC445333	-0.833215214
AI853383	TC417447	-0.831229999 RIKEN cDNA 2610002H19 gene (Mus musculus), complete
AU019316	TC399752	-0.830975347 homologue to sodium-phosphate cotransporter Ila C-terminal-associated protein 2 (Mus musculus), partial (46%)
BE380506	TC396816	-0.830730913 homologue to tricarboxylate carrier - rat (fragment), complete
AI845703	TC408471	-0.828943602 homologue to unknown (Homo sapiens), complete
AW206668	TC418017	-0.828941978 putative (Mus musculus), complete
AU015471	TC388035	-0.825695977 RIKEN cDNA 2610529C04 gene (Mus musculus), complete
AI853147	TC439869	-0.825658898 Unknown (protein for MGC:7461) (Mus musculus), complete
AI841995		-0.822490298
AI852401	TC368647	-0.81672979
AW536435	TC375933	-0.814783743 homologue to PROTEIN TRANSPORT PROTEIN SEC24D (SEC24-RELATED PROTEIN D). [Human], partial (53%)
AU016967	TC435441	-0.811845185 HIGH-AFFINITY COPPER UPTAKE PROTEIN 1 (HCTR1) (COPPER TRANSPORTER 1). [Human], partial (89%)
AI844113	TC374619	-0.810721367
AI845783	TC414247	-0.810640081
AI846919	TC407718	-0.809640474 glycerol-3-phosphate dehydrogenase (Mus musculus), complete
AI835613	TC385750	-0.805520877 putative (Mus musculus), complete
AW549443	TC410775	-0.8025098
AI847836	TC367785	-0.79883557 homologue to unnamed protein product (Homo sapiens), partial (90%)
AW545316	TC407919	-0.798255765
AA408946	TC407718	-0.797551213 glycerol-3-phosphate dehydrogenase (Mus musculus), complete
AI849518	TC375357	-0.797470116 homologue to putative (Mus musculus), complete
AI838680	TC396816	-0.797058635 homologue to tricarboxylate carrier - rat (fragment), complete
AA409879	TC393887	-0.79606833
AU016261	TC418109	-0.789157761
AU021857	TC390427	-0.788921632 SERINE PROTEASE PC6 PRECURSOR (EC 3.4.21.-) (SUBTILISIN/KEXIN-LIKE PROTEASE PC5) (CONVERTASE PC5). [Mouse], complete
AU022447	TC371165	-0.785898061
AI852646	TC421710	-0.779890994 RIKEN cDNA 1110014J22 gene (Mus musculus), complete
AA409749		-0.779878706
AU040150	TC421761	-0.777891872
X58149		-0.776795484
C79447		-0.77533826
AU022670	TC420805	-0.773590403

C79535	TC440104	-0.771861129	
A1844670	TC417436	-0.768596449	
A1843250	TC436166	-0.762666702	sialyltransferase (Mus musculus), complete
A1850874	TC367333	-0.76205266	ALDEHYDE DEHYDROGENASE, CYTOSOLIC (EC 1.2.1.3) (ALDH CLASS 1) (ALDHII) (ALDH-E1). [Mouse], complete
C80816	TC382510	-0.761269065	Similar to pleckstrin homology, Sec7 and coiled/coil domains, binding protein (Mus musculus), complete
AU022042	TC377104	-0.759400012	
A1848192	TC368327	-0.757104072	SOMATOSTATIN PRECURSOR [CONTAINS: ANTRIN; SOMATOSTATIN-28; SOMATOSTATIN-14]. [Mouse, Rat], complete
A1852659	TC368015	-0.75302365	similar to hypothetical protein KIAA0232 (Homo sapiens), partial (21%)
AU016653	TC413962	-0.752686866	Type II membrane protein of ER-mouse gene similar to alpha-mannosidase (Mus musculus), complete
A1848984	TC375251	-0.751141674	VACUOLAR PROTON TRANSLOCATING ATPASE 116 KDA SUBUNIT A ISOFORM 2 (V-ATPASE 116-KDA ISOFORM A2) (IMMUNE SUPPRESSOR FACTOR J6B7). [Mouse], partial (27%)
AW538597	TC421714	-0.745294054	similar to MONOCARBOXYLATE TRANSPORTER 7 (MCT 7) (MCT 6). [Human], complete
A1842290	TC421546	-0.744401799	hepatic transcription activator DBP - rat, complete
AU023610	TC446359	-0.744294396	
A1835654	TC414334	-0.740721826	putative (Mus musculus), partial (95%)
AU022245	TC374733	-0.740284883	homologue to TSC22-related inducible leucine zipper 1b (Mus musculus), partial (8%)
AW545573	TC417741	-0.740232071	CYTOKINE RECEPTOR COMMON GAMMA CHAIN PRECURSOR (GAMMA-C) (INTERLEUKIN-2 RECEPTOR GAMMA CHAIN) (IL-2R GAMMA CHAIN) (P64). [Mouse], complete
A1841892	TC379075	-0.739153817	putative (Mus musculus), complete
AU022383	TC424533	-0.738422126	
AW536069	TC404515	-0.737395723	RIKEN cDNA 0610006H10 gene (Mus musculus), complete
AW545455	TC421761	-0.736079727	
A1846656	TC440290	-0.734883514	
AW545954	TC414427	-0.73473901	homologue to Similar to PTD015 protein (Homo sapiens), complete
A1836143	TC414334	-0.734002094	putative (Mus musculus), partial (95%)
A1851829	TC444169	-0.729528188	similar to KIAA0779 protein (Homo sapiens), partial (93%)
A1835855	TC425033	-0.728853673	
A1836168	TC432717	-0.728353644	putative (Mus musculus), complete
A1851374	TC439727	-0.728352982	putative (Mus musculus), complete
A1842338	TC402496	-0.726367279	
A1842131	TC367600	-0.72339986	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1 (Mus musculus), complete
AW548139	TC389629	-0.718614351	homologue of S70447 F1Fo-ATP synthase complex Fo membrane domain f subunit (cattle, heart, mRNA, 403 nt) [Mus musculus], partial (94%)
AU016860	TC443955	-0.718199506	RAS-RELATED PROTEIN RAB-30. [Human], complete
A1847957	TC414441	-0.715156913	homologue to SEC63 protein (Homo sapiens), complete
A1851875	TC378420	-0.714319289	RIKEN cDNA 5730403B10 gene (Mus musculus), complete
A1846022	TC385796	-0.711957491	neurotactin (Mus musculus), complete
AW553129	TC403925	-0.710793692	Similar to Acetyl-CoA acyltransferase, 3-oxo acyl-CoA thiolase A, peroxisomal (Homo sapiens), complete
A1851735	TC378276	-0.710478402	
A1844586	TC396619	-0.710181685	
A1844295	TC414788	-0.708925135	PHOSPHATIDATE CYTIDYLYLTRANSFERASE 1 (EC 2.7.7.41) (CDP-DIGLYCERIDE SYNTHETASE 1) (CDP-DIGLYCERIDE PYROPHOSPHORYLASE 1) (CDP-DIACYLGLYCEROL SYNTHASE 1) (CDS 1) (CTP-PHOSPHATIDATE CYTIDYLYLTRANSFERASE 1) (CDP-DAG SYNTHETASE 1) (CDP-DG SYNTHETASE 1). [Rat], pa
A1853951	TC374880	-0.707255828	putative (Mus musculus), complete
AW553105	TC393978	-0.706478647	
A1850497	TC396754	-0.706354723	arachidonate 5-lipoxygenase - mouse (fragment), complete
A1836637	TC371758	-0.698721011	
A1836731		-0.694119291	
A1853800	TC444203	-0.692350167	inner mitochondrial membrane peptidase 2 (Mus musculus), complete
A1844331	TC378407	-0.68904712	DELTA 1-PYRROLINE-5-CARBOXYLATE SYNTHETASE (P5CS) [INCLUDES: GLUTAMATE 5-KINASE (EC 2.7.2.11) (GAMMA-GLUTAMYL KINASE) (GK); GAMMA-GLUTAMYL PHOSPHATE REDUCTASE (GPR) (EC 1.2.1.41) (GLUTAMATE-5-SEMIALDEHYDE DEHYDROGENASE) (GLUTAMYL-GAMMA-SEMIALDEHYDE DE
AU043215	TC417933	-0.686314522	G21 protein (Mus musculus), complete
A1844065	TC405909	-0.683703152	
A1835105	TC396661	-0.68285918	monoglyceride lipase (Mus musculus), complete
A1119976	TC422231	-0.682294532	transforming protein (cbl) - mouse, complete
A1848335	TC425422	-0.674961455	similar to Similar to outer dense fiber of sperm tails 2 (Homo sapiens), partial (12%)
AU017550	TC411387	-0.672461351	
A1841894	TC396919	-0.669710709	putative (Mus musculus), partial (62%)
A1838190	TC415090	-0.665081067	
A1848783	TC444383	-0.664830915	[PYRUVATE DEHYDROGENASE [LIPOAMIDE]] KINASE ISOZYME 2, MITOCHONDRIAL PRECURSOR (EC 2.7.1.99) (PYRUVATE DEHYDROGENASE KINASE ISOFORM 2). [Mouse], complete
A1853999	TC443215	-0.6623952	homologue to RAS-RELATED PROTEIN RAB-9 (ONCOGENE C-MEL). [Human, Dog], complete
C87042	TC391601	-0.659570939	
A1839956	TC378890	-0.658431582	
A1839625	TC402399	-0.658363283	
AU017653	TC408748	-0.656994881	homologue to hypothetical protein (Macaca fascicularis), partial (36%)
C85175	TC417527	-0.652677066	homologue to putative (Mus musculus), partial (12%)
A1850982	TC414932	-0.652606872	putative (Mus musculus), complete
AU041795	TC409302	-0.647693568	mDab555 protein (Mus musculus), complete
A1854136	TC411211	-0.646255125	
A1846408	TC433177	-0.638065653	
A1854218	TC393533	-0.637890563	doublecortin-like kinase (Mus musculus), complete
A1838566	TC404719	-0.634900347	structural protein FBF1 (Mus musculus), complete
A1846925	TC375465	-0.633469441	homologue to LONG-CHAIN-FATTY-ACID-COA LIGASE, BRAIN ISOZYME (EC 6.2.1.3) (LONG-CHAIN ACYL-COA SYNTHETASE) (LACS). [Rat], partial (72%)
A1853433		-0.633230428	
A1847702	TC386329	-0.632242659	apoptosis activator Mtd; Bcl-2-related ovarian killer protein-like; apoptosis activator [Mus musculus]
A1840624	TC400957	-0.629934506	putative (Mus musculus), complete
A1852213	TC397001	-0.62929356	UDP-GLUCOSE 6-DEHYDROGENASE (EC 1.1.1.22) (UDP-GLC DEHYDROGENASE) (UDP-GLCDH) (UDPGDH). [Mouse], complete
AU045737	TC389972	-0.62897852	homologue to Nadrin1 (Mus musculus), complete
AU045930	TC417959	-0.628264553	homologue to transcription regulator - mouse, complete
A1851636	TC371886	-0.626531832	FORMIN 2. [Mouse], complete
C81014		-0.626158467	
A1853902	TC414806	-0.623185523	SCN8A (Mus musculus), complete
C85252	TC432571	-0.622714975	homologue to chromogranin/secretogranin-like vesicle protein precursor - rat, partial (75%)
A1853944	TC379054	-0.62173456	Similar to signaling intermediate in Toll pathway-evolutionarily conserved (Mus musculus), complete
AU023733	TC394697	-0.62088977	weakly similar to ribonuclease/angiogenin inhibitor 2 (Mus musculus), partial (29%)
A1852719	TC371026	-0.620884275	
AU024360	TC391903	-0.619902482	
A1838914	TC391777	-0.61955321	homologue to ankryrin repeat-rich membrane-spanning protein (Rattus norvegicus), partial (37%)
AU040320	TC435997	-0.618901315	homologue to unnamed protein product (Homo sapiens), partial (87%)
AW558117	TC437292	-0.617244952	
A1851597	TC371723	-0.614780485	ELONGATION FACTOR 1-ALPHA 2 (EF1-ALPHA-2) (ELONGATION FACTOR 1 A-2) (EEF1A-2) (STATIN S1). [Mouse, Rat], complete
A1835632	TC417673	-0.614550956	similar to KIAA1042 protein (Homo sapiens), partial (23%)
A1841151	TC429438	-0.613660913	DM-20 (Homo sapiens), complete
AU022330	TC410374	-0.612744859	
A1840716		-0.611657936	
A1847088	TC404197	-0.608315552	sialic acid-specific 9-O-acetyltransferase (Mus musculus), complete
AU021686	TC432961	-0.605360251	
AU014892	TC371462	-0.600554446	

AIB44651 TC374953 -0.599385364 similar to ZINC FINGER PROTEIN 183, [Human], complete
C85795 TC390192 -0.594755393 homologue to unnamed protein product (Homo sapiens), complete
AA07369 TC362438 -0.592432783 Similar to NADH dehydrogenase (ubiquinone) Fe-S protein 5 (15kD) (NADH-coenzyme Q reductase) (Mus musculus), complete
AW55655 TC390322 -0.59229730
C86619 TC367955 -0.574923651 similar to tetratricopeptide repeat domain 4 (Homo sapiens), complete
AIB48043 TC375699 -0.571334537 putative (Mus musculus), complete
AW548274 TC431411 -0.550162946
AIB49568 TC393239 0.558960005
AW536088 TC435718 0.560222342 ribonucleoside-diphosphate reductase (EC 1.17.4.1) chain M1 - mouse, complete
AUI016390 TC414169 0.570812117 GTP-binding protein NGB (Mus musculus), partial (27%)
AIB37094 TC394198 0.575056934 Similar to neurexophilin 3 (Mus musculus), complete
AW554829 TC422182 0.576151678 similar to bA305P22.2.1 (novel protein, isoform 1) (Homo sapiens), complete
AW554530 TC385785 0.576776499 PROCOLLAGEN C-PROTEINASE ENHANCER PROTEIN PRECURSOR (PCPE) (TYPE I PROCOLLAGEN COOH-TERMINAL PROTEINASE ENHANCER) (TYPE 1 PROCOLLAGEN C- PROTEINASE ENHANCER PROTEIN) (P14). [Mouse], complete
AW554403 0.577923929
AIB43647 TC414080 0.577963396 homologue to probable transcription factor CA150 - human, complete
AIB41098 TC376138 0.582858864
AIB47285 TC388634 0.589505701 apoptosis signal-regulating kinase 2; mitogen-activated protein kinase kinase 6; apoptosis signal-regulating kinase 2 [Mus musculus];
AIB54439 TC401164 0.592088142
AW554225 TC395280 0.598741424
AIB42636 TC382540 0.599557961 Similar to phenylalanine-tRNA synthetase (Mus musculus), complete
AU043814 TC435055 0.600971098
AW549430 TC400794 0.601158361 putative (Mus musculus), complete
AIB53274 TC403902 0.601710145 SPHINGOMYELIN PHOSPHODIESTERASE PRECURSOR (EC 3.1.4.12) (ACID SPHINGOMYELINASE). [Mouse], complete
AIB53829 TC414725 0.606560766
AW538624 TC403956 0.608056791 Similar to polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I) (Mus musculus), partial (21%)
AU044860 TC371736 0.608772201
AW537613 TC378301 0.608953639 NDRG1 PROTEIN (N-MYC DOWNSTREAM REGULATED GENE 1 PROTEIN) (NDRG1). [Mouse], complete
AIB48890 TC404846 0.611227495 GROWTH ARREST AND DNA-DAMAGE-INDUCIBLE PROTEIN GADD45 BETA (NEGATIVE GROWTH-REGULATORY PROTEIN MYD118) (MYELOID DIFFERENTIATION PRIMARY RESPONSE PROTEIN MYD118). [Mouse], complete
AW543707 TC393520 0.611582833 RIKEN cDNA 5730466A07 gene (Mus musculus), complete
AW553290 0.611885013
AW548569 TC400780 0.612141138 putative (Mus musculus), complete
AU017373 TC418169 0.612364687 putative (Mus musculus), complete
AIB53508 TC417591 0.614661291
C80533 0.619979999
AW537825 TC393509 0.622892854 similar to RES4-25 protein - human, partial (71%)
AW551281 TC372039 0.624738525
AIB36378 TC382648 0.625239231
AW553404 TC436089 0.625414544 homologue to ORIGIN RECOGNITION COMPLEX SUBUNIT 6. [Mouse], complete
AIB49000 TC389626 0.625846062 MATRIX METALLOPROTEINASE-14 PRECURSOR (EC 3.4.24.-) (MMP-14) (MEMBRANE-TYPE MATRIX METALLOPROTEINASE) (MT1-MMP) (MT1MMP) (MMP-X1) (MT-MMP). [Mouse], complete
AIB53099 TC388886 0.627485254
AIB47226 TC437811 0.628024304 homologue to small rec (Rattus norvegicus), partial (25%)
AW542466 TC407573 0.629732872 polypeptide GalNAc transferase-T2 (Mus musculus), complete
AU016751 TC393419 0.631284411 homologue to S164 (Homo sapiens), partial (74%)
AW538403 TC435846 0.631293433 formin binding protein 11 (Mus musculus), complete
AIB39000 TC382107 0.632087589 putative (Mus musculus), complete
AW552373 TC436416 0.633517526 CASPASE-12 PRECURSOR (EC 3.4.22.-). [Mouse], complete
AW541463 TC429591 0.636431646 putative (Mus musculus), complete
AIB47129 TC432947 0.638015763 CORTICOSTEROID 11-BETA-DEHYDROGENASE, ISOZYME 1 (EC 1.1.1.146) (11-DH) (11-BETA-HYDROXYSTEROID DEHYDROGENASE 1) (11-BETA-HSD1) (11BETA-HSD1A). [Mouse], complete
AIB47148 TC425322 0.638986285 cadherin-5 (Mus musculus), complete
AU023036 TC387995 0.640416174
AW536111 TC428915 0.645585835 putative (Mus musculus), complete
AIB45864 0.646728201
AIB46427 TC414872 0.647751825
AW553876 TC382483 0.647840121 similar to KIAA1025 protein (Homo sapiens), partial (75%)
AIB48763 TC375650 0.647945827 OL-protocadherin isoform (Mus musculus), partial (40%)
AU041258 TC439751 0.649264102 ethanol induced gene product (Mus musculus), complete
AIB49312 TC421481 0.650717901 JUNCTIONAL ADHESION MOLECULE PRECURSOR (JAM). [Mouse], complete
AW548009 TC424840 0.65180625 homologue to putative (Mus musculus), complete
AIB36955 TC419059 0.653040399 homologue to KIAA1235 protein (Homo sapiens), partial (33%)
AUI019831 TC393419 0.654069826 homologue to S164 (Homo sapiens), partial (74%)
AU023931 TC394828 0.654843308 similar to CARBONIC ANHYDRASE IX PRECURSOR (EC 4.2.1.1) (CARBONATE DEHYDRATASE IX) (CA-X) (CAIX) (MEMBRANE ANTIGEN MN) (P54/58N) (RENAL CELL CARCINOMA-ASSOCIATED ANTIGEN G250) (RCC-ASSOCIATED ANTIGEN G250) (PMW1). [Human], partial (23%)
AW539587 TC436131 0.655165171 CMP-SIALIC ACID TRANSPORTER (CMP-SIA-TR) (CMP-SA-TR). [Mouse], complete
AW552014 TC386261 0.657308128 transcriptional repressor SIN3B (Mus musculus), complete
AU017699 TC371354 0.65869738 similar to unnamed protein product (Homo sapiens), partial (38%)
AW542918 TC371870 0.659177146 Similar to beta-glucuronidase structural (Mus musculus), complete
C81271 TC417359 0.661849851 putative (Mus musculus), complete
AIB37483 TC388510 0.662194997
AIB45860 TC389864 0.663530352 CTLA-2-ALPHA PROTEIN PRECURSOR. [Mouse], complete
AW536539 TC439789 0.667190329 homologue to AMIDOPHOSPHORIBOSYLTRANSFERASE PRECURSOR (EC 2.4.2.14) (GLUTAMINE PHOSPHORIBOSYL-PYROPHOSPHATE AMIDOTRANSFERASE) (ATASE) (GPAT). [Rat], partial (38%)
AW537769 TC382228 0.668659193 putative (Mus musculus), complete
AIB39527 TC379690 0.669628638 homologue to AMYLOID BETA A4 PRECURSOR PROTEIN-BINDING FAMILY A MEMBER 1 (NEURON-SPECIFIC X11 PROTEIN) (NEURONAL MUNC18-1-INTERACTING PROTEIN 1) (MINT-1) (ADAPTER PROTEIN X11ALPHA). [Rat], partial (23%)
AW536344 TC425049 0.671665808 homologue to unnamed protein product (Homo sapiens), complete
AW543975 TC375371 0.673831249
AW552186 TC424685 0.674373596 Similar to ubiquitin carrier protein (Mus musculus), complete
AIB39009 TC403923 0.675451499 Unknown (protein for MGC:12123) (Mus musculus), complete
AW547303 TC385711 0.675467397 NEDD-4 PROTEIN (EC 6.3.2.-) (FRAGMENT). [Mouse], partial (11%)
AW554923 TC428407 0.678261899 putative (Mus musculus), complete
AIB39441 TC393687 0.679508474 protein kinase C and casein kinase II substrate 3 (Mus musculus), complete
AUI041740 TC443079 0.680995784 FIBRINOMODULIN PRECURSOR (FII). [Mouse], complete
AW553387 TC383763 0.680258689 similar to unnamed protein product (Homo sapiens), partial (17%)
AU017659 TC397068 0.682565463 EH domain protein Reps1 - mouse, complete
AU044053 TC371543 0.688943939 g1-related zinc finger protein (Mus musculus), complete
AIB41886 TC381986 0.689364742 procollagen, type I, alpha 2 (Mus musculus), complete
AIB38797 TC432397 0.693139704 putative (Mus musculus), complete
AW556448 TC398690 0.693211905 putative (Mus musculus), complete
AU016519 TC407761 0.693310673 putative (Mus musculus), complete
AW552870 TC397130 0.693798955 homologue to hypothetical protein (Macaca fascicularis), partial (81%)
AW557980 TC414645 0.696532898 [PYRUVATE DEHYDROGENASE [LIPOAMIDE]]-PHOSPHATASE 1, MITOCHONDRIAL PRECURSOR (EC 3.1.3.43) (PDP 1) (PYRUVATE DEHYDROGENASE PHOSPHATASE, CATALYTIC SUBUNIT 1) (PPDC 1). [Rat], partial (44%)
AW537736 TC433008 0.697547864 putative (Mus musculus), complete
AW556761 TC375156 0.698579249 putative (Mus musculus), complete
BES31475 TC386211 0.699810407 neuronal differentiation related protein (Mus musculus), complete
AIB52142 TC435987 0.70264976 myosin I alpha chain - mouse, partial (22%)

AI848244	TC439828	0.705130098 putative (Mus musculus), partial (95%)
AI852481	TC440214	0.705612793 L-MYC PROTO-ONCOGENE PROTEIN [Mouse], complete
AI852760	TC432149	0.709900852
AI853345	TC421303	0.711125832 Similar to dimethylarginine dimethylaminohydrolase 2 (Mus musculus), complete
AI850137	TC378285	0.711876992 nucleolin - mouse, complete
AI853111	TC429442	0.713580143 putative (Mus musculus), complete
AU022931	TC371388	0.716529463 homologue to prostein (Homo sapiens), complete
AW551740	TC443869	0.718078657 putative (Mus musculus), complete
AI848817	TC394370	0.721947075 4-1BB LIGAND (4-1BBL) [Mouse], complete
AW551445	TC414715	0.722426627
AI850456	TC393309	0.72292725 NAD(P)-DEPENDENT STEROID DEHYDROGENASE (EC 1.1.1.-) [Mouse], complete
AW538481	TC382094	0.723328145 Similar to eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD) (Mus musculus), complete
AI851060	TC439588	0.726460367 beta-glucuronidase (EC 3.2.1.31) allele B precursor - mouse, complete
AU043264	TC443887	0.73105855 transcription repressor CRTR-1 (Mus musculus), complete
AI846628	TC367931	0.734470516 homologue to anaphase-promoting complex subunit 4 (Homo sapiens), partial (97%)
AW536141	TC392521	0.734785255 putative (Mus musculus), complete
AI835583	TC443151	0.737668008 glutamate-ammonia ligase (Mus musculus), complete
AU040277	TC403923	0.738975379 Unknown (protein for MGC:12123) (Mus musculus), complete
AW549534	TC426472	0.741964489
AI853298	TC407788	0.747326106 transcription activator SRG3 - mouse, complete
AW557035	TC386514	0.748587695 putative (Mus musculus), complete
AU017975	TC439607	0.750315828 homologue to EH domain binding protein epsin 2 (Rattus norvegicus), complete
AW553970	TC463961	0.752816526 putative (Mus musculus), complete
AU018183	TC425220	0.75329461 similar to Similar to hypothetical protein FLJ14075 (Homo sapiens), partial (55%)
AW556921	TC375070	0.753447507 similar to melanoma-associated antigen MG50 (Homo sapiens), partial (42%)
AU045867		0.754504201
AI836854	TC417359	0.755216531 putative (Mus musculus), complete
AU047111	TC445294	0.755942542
AU018592	TC409534	0.755947823
C76390	TC389758	0.756065552 eIF-1A (Mus musculus), complete
C87372	TC375511	0.757323165 KIAA1771 protein (Homo sapiens), partial (16%)
AI848139	TC413702	0.758217808
AW536886	TC428434	0.760896096 putative (Mus musculus), complete
AW547245		0.76165005
AW546835	TC432864	0.762923848 homologue to putative (Mus musculus), complete
AI838819	TC382172	0.763604101 myb-binding protein p160 - mouse, complete
AI835674	TC443151	0.76371302 glutamate-ammonia ligase (Mus musculus), complete
AW554486	TC397249	0.764823123 Similar to ubiquitous nuclear protein (Mus musculus), complete
AI846046	TC404104	0.767579326 Acyl-CoA synthetase, long chain (Mus musculus), complete
AW535927	TC439528	0.768453053 homologue to MEG3 (Homo sapiens), complete
C77170	TC417361	0.768599476
AW540967	TC389542	0.768762176 unnamed protein product (Homo sapiens), complete
AW536780	TC407788	0.769291345 transcription activator SRG3 - mouse, complete
AI845405	TC389849	0.76939595 putative (Mus musculus), complete
AA409082	TC443213	0.770173727 homologue to talpoxin-associated calcium binding protein 49 (Mus musculus), complete
AW553551	TC404449	0.770703926 homologue to CL2B2 protein - rat, partial (29%)
AW539467	TC428189	0.771239606 poly A binding protein, cytoplasmic 1 (Mus musculus), complete
AI848400	TC407737	0.778870129
C78116	TC432570	0.779666968 similar to unnamed protein product (Homo sapiens), partial (96%)
AW548068	TC435472	0.780428071 epsilon-globin (Mus musculus), complete
AW550265	TC414087	0.783093203 Unknown (protein for IMAGE:3487926) (Mus musculus), partial (97%)
AI847971	TC374591	0.792724202 putative (Mus musculus), complete
AW539335	TC404992	0.792916292 homologue to KIAA0532 protein (Homo sapiens), partial (18%)
AA408450	TC399637	0.793516276 homologue to STRUCTURE-SPECIFIC RECOGNITION PROTEIN 1 (SSRP1) (RECOMBINATION SIGNAL SEQUENCE RECOGNITION PROTEIN) (T160) [Mouse], partial (31%)
AW551658	TC367657	0.795866275 putative (Mus musculus), complete
AU042966	TC368460	0.800625636 similar to unnamed protein product (Homo sapiens), partial (32%)
AW543081	TC425242	0.802894653 uridine-cytidine kinase 2 (Mus musculus), complete
AW551943	TC390446	0.805336395
AI847288	TC390792	0.806846987 putative (Mus musculus), complete
AW543698	TC425322	0.810681786 cadherin-5 (Mus musculus), complete
AI841827	TC386316	0.814612531 putative (Mus musculus), complete
AW536751	TC378632	0.815841958 SERINE-THREONINE-PROTEIN KINASE PLK (EC 2.7.1.-) (PLK-1) (SERINE-THREONINE PROTEIN KINASE 13) (STPK13) [Mouse], complete
AI845798	TC379205	0.819672389 putative (Mus musculus), complete
AW548769	TC436039	0.826675601 putative (Mus musculus), complete
AW545839	TC435582	0.82791425 NUCLEOSOME ASSEMBLY PROTEIN 1-LIKE 1 (NAP-1 RELATED PROTEIN) (BRAIN PROTEIN DN38) [Mouse], partial (82%)
AU045027	TC375455	0.828444151
AW552829	TC376999	0.828843846
AI841088	TC414531	0.830329899
AI842868	TC390022	0.830779868 homologue to adapter protein RUK-1 (Rattus norvegicus), partial (91%)
AW549810	TC404205	0.831651465 semaF cytoplasmic domain associated protein 1 (Mus musculus), complete
AI835221	TC400510	0.837654184 putative (Mus musculus), complete
AI842576	TC421401	0.83845666 potassium channel, subfamily K, member 1 (Mus musculus), complete
AW537169	TC371902	0.838896582 Similar to hypothetical protein FLJ10504 (Mus musculus), complete
AW541452	TC371434	0.839270357 (RNA-guanine transglycosylase (Mus musculus), complete
AU042961	TC397148	0.841213915 collagen alpha 1(XVII) chain precursor, long splice form - mouse, complete
AA410041	TC425243	0.842046995 putative (Mus musculus), complete
AU021260	TC422063	0.843920797 similar to unnamed protein product (Homo sapiens), partial (31%)
AU021357	TC393690	0.846367039 KIAA1063 protein (Homo sapiens), partial (55%)
AI849357	TC387038	0.847135029 PRO-NEUREGULIN-3 PRECURSOR (PRO-NRG3) [CONTAINS: NEUREGULIN-3 (NRG-3)] [Mouse], complete
AU041966	TC436719	0.847190358 MINIMUM POTASSIUM ION CHANNEL-RELATED PEPTIDE 2 (MIRP2) (MINK-RELATED PEPTIDE 2) [Mouse], complete
C79711		0.850418244
AW547996	TC400736	0.850505827 similar to unnamed protein product (Homo sapiens), complete
AW547850	TC443116	0.858244961 protocadherin gamma A1 (Mus musculus), complete
AI848921	TC384927	0.85927846
AI841664	TC435583	0.86370427 homologue to putative (Mus musculus), partial (37%)
AW550056	TC390309	0.867124807 putative (Mus musculus), complete
AU017612		0.868973644
AI843056	TC415159	0.872701351 Cegp1 protein (Mus musculus), complete
AW536858	TC428709	0.873046141 similar to This sequence is almost identical to HUMRSC433, D13632. (Homo sapiens), complete
AW554413	TC441833	0.876026263
AW536278	TC440214	0.877729969 L-MYC PROTO-ONCOGENE PROTEIN [Mouse], complete
AI846565	TC417422	0.880791221 similar to P63 protein - human, complete

AW545287 TC414899 0.881495752
AB35858 TC432201 0.882372892 homologue to TROPOMYOSIN, FIBROBLAST NON-MUSCLE TYPE (TROPOMYOSIN 4) (TM30-PL). [Human, Pig], complete
CB0159 TC374864 0.882647968 ISOPENTENYL-DIPHOSPHATE DELTA-ISOMERASE (EC 5.3.3.2) (IPP ISOMERASE) (ISOPENTENYL PYROPHOSPHATE ISOMERASE). [Mouse], complete
BE329376 TC390346 0.883183472 mage-d1 (Mus musculus), complete
AW538689 TC389693 0.885133187 similar to myelin protein zero-like 1 (Homo sapiens), complete
AW553366 TC374736 0.885296895 putative (Mus musculus), complete
AW548804 TC424982 0.885780022 In1b (Homo sapiens), partial (70%)
C75777 TC376771 0.886978982 putative (Mus musculus), complete
AW549993 TC404354 0.887070659 weakly similar to alcohol dehydrogenase Pan1b (Mus musculus), partial (66%)
AB47448 TC372332 0.88707169
AW553381 TC438233 0.887483682
AB151214 TC404053 0.891111175 putative (Mus musculus), complete
AW559158 TC443026 0.891786743 COLLAGEN ALPHA 1 (IV) CHAIN PRECURSOR. [Mouse], complete
AB42165 TC421224 0.892017319 GLUTATHIONE S-TRANSFERASE MU 5 (EC 2.5.1.18) (GSTM5) (GST CLASS-MU) (FIBROUS SHEATH COMPONENT 2) (FSC2). [Mouse], complete
AW536420 TC413757 0.894732303 6-PHOSPHOFRUCTOKINASE, LIVER TYPE (EC 2.7.1.11) (PHOSPHOFRUCTOKINASE 1) (PHOSPHOHEXOKINASE) (PHOSPHOFRUCTO-1-KINASE ISOZYME B) (PFK-B). [Mouse], complete
AB45434 TC379256 0.897920091 THROMBOSPONDIN 2 PRECURSOR. [Mouse], complete
AW546280 TC407735 0.904489021
AU021239 TC425131 0.906674984 homologue to Clast4 protein (Mus musculus), complete
AW744595 TC394579 0.917741382 MACROPHAGE SCAVENGER RECEPTOR TYPES I AND II (MACROPHAGE ACETYLATED LDL RECEPTOR I AND II) (SCAVENGER RECEPTOR TYPE A) (SR-A). [Mouse], complete
AW546921 TC382334 0.921047104 LYMPHOCTE ANTIGEN LY-6A.2/LY-6E.1 PRECURSOR (T-CELL-ACTIVATING PROTEIN) (TAP). [Mouse], complete
AW551726 TC378584 0.923043519 WW domain binding protein 5 (Mus musculus), complete
AW537992 TC418006 0.923492343 putative (Mus musculus), complete
AW637045 TC404040 0.923612245 METHIONINE AMINOPEPTIDASE 2 (EC 3.4.11.18) (METAP 2) (PEPTIDASE M 2) (INITIATION FACTOR 2 ASSOCIATED 67 KDA GLYCOPROTEIN) (P67). [Mouse], complete
AU041881 TC382323 0.924998625 TAJ-alpha long (Mus musculus), complete
AW556448 TC345841 0.933458759 DICKKOPF RELATED PROTEIN-3 PRECURSOR (DKK-3) (DICKKOPF-3) (MDKK-3). [Mouse], complete
C76763 TC424920 0.935375245 similar to unnamed protein product (Homo sapiens), complete
AB42533 TC431995 0.936246313 macrophage migration inhibitory factor DER6 - mouse, complete
C77965 TC414312 0.938402547 aldose reductase homolog - mouse, complete
AB430335 TC429237 0.938654574 EXCITATORY AMINO ACID TRANSPORTER 2 (SODIUM-DEPENDENT GLUTAMATE/ASPARTATE TRANSPORTER 2). [Mouse], complete
AW536773 TC400510 0.942078602 putative (Mus musculus), complete
AW549121 TC417392 0.943033054 nonhistone chromosomal protein HMG-14 - mouse, complete
AW557319 TC375446 0.949278585 putative (Mus musculus), complete
AW541476 TC390031 0.952176872
AW546616 TC386066 0.952686246 Similar to interferon induced transmembrane protein 3 (1-8U) (Mus musculus), complete
CB6354 TC435548 0.958528672 serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 2 (Mus musculus), complete
AB42540 TC385583 0.960721409 NC1 protein (Rattus norvegicus), complete
AU042596 TC371633 0.96100507 unnamed protein product (Homo sapiens), partial (86%)
AB151618 TC436370 0.961383885 homologue to Similar to sex comb on midleg-like 1 (Drosophila) (Homo sapiens), complete
AB39275 TC443026 0.962227678 COLLAGEN ALPHA 1 (IV) CHAIN PRECURSOR. [Mouse], complete
AW554493 TC441452 0.963393769
AW546767 0.964531776
AB40441 TC432571 0.965708213 homologue to chromogranin/secretogranin-like vesicle protein precursor - rat, partial (75%)
AW551743 TC415576 0.968127082
AB353088 TC413774 0.9702407 follistatin-like (Mus musculus), complete
AW539558 TC439940 0.970786446
AW554737 TC443615 0.97256332 homologue to putative (Mus musculus), partial (85%)
AW545351 TC417471 0.976246922 Unknown (protein for MGC:6446) (Mus musculus), complete
AW544184 0.980307334
AB38277 TC414417 0.98204968 ETS-DOMAIN PROTEIN ELK-3 (ETS-RELATED PROTEIN NET) (ETS-RELATED PROTEIN ERP). [Mouse], complete
AW550920 TC390133 0.985015516 vascular endothelial zinc finger 1 (Mus musculus), partial (27%)
AB353558 TC400881 0.985955914 type I transmembrane protein Fn14 (Mus musculus), complete
AW543423 TC378571 0.98923873 glucose transport protein GT1 - mouse, complete
AW544680 TC405365 0.990486319 putative (Mus musculus), complete
BE31947 TC378258 0.991297439 milk fat globule glycoprotein MFG-E8 S (Mus musculus), complete
BE457675 TC431995 0.992579858 macrophage migration inhibitory factor DER6 - mouse, complete
AW210164 TC431952 0.993675425 CYCLIN-DEPENDENT KINASE INHIBITOR 1C (CYCLIN-DEPENDENT KINASE INHIBITOR P57) (P57KIP2). [Mouse], partial (97%)
AB46883 TC375038 1.000170595 RIKEN cDNA 2310056P07 gene (Mus musculus), complete
AW536614 TC368093 1.004696351
AB39424 TC418185 1.005919677
AW555131 1.008188337
AU017967 TC429536 1.01372296 MATRILYSIN PRECURSOR (EC 3.4.24.23) (PUMP-1 PROTEASE) (UTERINE METALLOPROTEINASE) (MATRIX METALLOPROTEINASE-7) (MMP-7) (MATRIN). [Mouse], complete
AW229040 TC385801 1.014337874 PROTEIN-LYSINE 6-OXIDASE PRECURSOR (EC 1.4.3.13) (LYSYL OXIDASE) (RAS EXCISION PROTEIN). [Mouse], complete
AW547664 TC397557 1.015994838
AW547573 TC386806 1.016779103 weakly similar to dJ329L24.1 (novel protein) (Homo sapiens), partial (96%)
AB44467 1.016805703
AB353554 TC371949 1.029112513 homologue to multidrug resistance protein (Mus musculus), partial (15%)
AW544893 TC428557 1.035963399 putative (Mus musculus), complete
AU015495 TC432254 1.036748273 putative (Mus musculus), complete
AB41954 TC428544 1.037133674
AW546291 TC432828 1.038845297
AW551393 TC436345 1.039877819 TRANSCRIPTION FACTOR E2F6 (E2F-6) (E2F-BINDING SITE MODULATING ACTIVITY PROTEIN) (EMA). [Mouse], complete
AB41796 TC371998 1.040400783
AU022324 TC443373 1.042949488 legumain (Mus musculus), complete
AU045060 TC421743 1.044101285 similar to Ig gamma-2b chain - mouse, complete
AU019064 TC372235 1.046173851 ets transcription factor - mouse, complete
AU016757 TC404271 1.049265705 MYC PROTO-ONCOGENE PROTEIN (C-MYC). [Mouse], complete
AU017822 TC421114 1.061881585
AB39543 TC397857 1.062603625 VOLTAGE-DEPENDENT P/Q-TYPE CALCIUM CHANNEL ALPHA-1A SUBUNIT (CALCIUM CHANNEL, L TYPE, ALPHA-1 POLYPEPTIDE ISOFORM 4) (BRAIN CALCIUM CHANNEL I) (BI). [Mouse], complete
AW544543 TC431995 1.064076493 macrophage migration inhibitory factor DER6 - mouse, complete
AB52349 TC368207 1.064407796 Similar to hypothetical protein (Mus musculus), complete
AB38745 1.065575855
AW542320 TC387590 1.071012632
AW536425 TC378285 1.074931129 nucleolin - mouse, complete
AU040803 TC432752 1.07529553 calcium binding protein D-9k; CabP9k (Mus musculus), complete
AB46486 TC378337 1.076409425 homologue to Unknown (protein for MGC:9749) (Homo sapiens), complete
AU040767 TC431956 1.07728768 p57KIP2 (Mus musculus), partial (49%)
AW541453 TC407678 1.079160557 Similar to capping protein (actin filament), gelsolin-like (Mus musculus), complete
AW546463 TC400713 1.081027332 PLM-like protein (Mus musculus), complete
AW557617 TC432373 1.082736349 putative (Mus musculus), complete
AW544811 TC435654 1.086243877 METALLOTHIONEIN-I (MT-I). [Mouse], complete
AW555633 TC385243 1.089001668
AB42770 TC439488 1.091960801 plastin 3 (T isoform) (Mus musculus), complete

AW549841	TC371319	1.093617825 putative (Mus musculus), complete
AW543683	TC378285	1.096569717 nucleolin - mouse, complete
AW549877	TC425246	1.098178222 homologue to KIAA1722 protein (Homo sapiens), partial (11%)
AI849247	TC374984	1.11729552 E1B 19k/Bcl-2-binding protein homolog putative (Mus musculus)
AW538700	TC414563	1.102029905 similar to unnamed protein product (Homo sapiens), partial (37%)
AU018106	TC420527	1.105445007
AW549829	TC389371	1.106537756
AW549872	TC383030	1.10798273 homologue to PROTEIN-TYROSINE PHOSPHATASE GAMMA PRECURSOR (EC 3.1.3.48) (R-PTP- GAMMA), [Mouse], complete
AW553526	TC385716	1.111880919 NUCLEOPHOSMIN (NPM) (NUCLEOLAR PHOSPHOPROTEIN B23) (NUMATRIN) (NUCLEOLAR PROTEIN NO38), [Mouse], complete
AW553554	TC387803	1.11729552 E1B 19k/Bcl-2-binding protein homolog putative (Mus musculus)
AI854178	TC397452	1.117652481 lanosterol 14-alpha-demethylase (Mus musculus), complete
AU042018	TC393273	1.117870462 putative (Mus musculus), complete
AI837648	TC408381	1.120214459
AI836247		1.122502234
AW541474	TC378285	1.122524127 nucleolin - mouse, complete
AI836644	TC379167	1.123802833 homologue to KIAA1749 protein (Homo sapiens), partial (38%)
BE381030	TC421004	1.124763776 COLLAGEN ALPHA 1(I) CHAIN PRECURSOR, [Mouse], complete
BE532311	TC418200	1.127649668 hypoxia-inducible factor 1 alpha (Mus musculus), complete
AW553526	TC385716	1.128320626 NUCLEOPHOSMIN (NPM) (NUCLEOLAR PHOSPHOPROTEIN B23) (NUMATRIN) (NUCLEOLAR PROTEIN NO38), [Mouse], complete
AI842716	TC378578	1.139393713
AI854349	TC424781	1.140663337 SERINE/THREONINE-PROTEIN KINASE SGK (EC 2.7.1.-) (SERUM/GLUCOCORTICOID-REGULATED KINASE), [Mouse], complete
CB8016	TC443151	1.14761282 glutamate-ammonia ligase (Mus musculus), complete
AW554924	TC371272	1.147640031 position-effect variegation 3-9 homolog (Mus musculus), partial (96%)
AI837598		1.148570709
C76481	TC394108	1.150184731 urokinase-type plasminogen activator receptor, hypothetical soluble form precursor - mouse, complete
AI854495	TC439713	1.150521301 Similar to glycogenin 2 (Mus musculus), complete
AI853660	TC429088	1.15100197 LAMININ ALPHA-5 CHAIN (FRAGMENT), [Mouse], complete
AW545145	TC439313	1.155923993 homologue to pantophysin (Mus musculus), partial (95%)
AW545796	TC415542	1.155908467
AI852411	TC400479	1.172846062 homologue to Unknown (protein for IMAGE:3590584) (Mus musculus), complete
AI841112	TC371355	1.172888786 putative (Mus musculus), complete
AU018274	TC431975	1.177649566 CALPAIN 2, LARGE [CATALYTIC] SUBUNIT (EC 3.4.22.17) (CALCIUM-ACTIVATED NEUTRAL PROTEINASE) (CANP) (M-TYPE) (MILLIMOLAR-CALPAIN) (80 KDA M- CALPAIN SUBUNIT) (CALP80), [Mouse], complete
AI846732	TC368174	1.1802162 Unknown (protein for MGC:13786) (Mus musculus), complete
AW537036	TC379096	1.182673289 duit1 protein - mouse, complete
AI841326	TC435418	1.190423463 Similar to Transferrin (Mus musculus), complete
AI844072	TC435848	1.210333376 serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 2 (Mus musculus), complete
AU040478	TC445273	1.218011614
AU040774		1.220844398
AW548203	TC404840	1.222346007 putative (Mus musculus), complete
AW554541		1.233707355
AU040460	TC383173	1.234616361 putative (Mus musculus), complete
AW208825	TC432447	1.26204854 CORTICOSTEROID 11-BETA-DEHYDROGENASE, ISOZYME 1 (EC 1.1.1.146) (11-DH) (11-BETA-HYDROXYSTEROID DEHYDROGENASE 1) (11-BETA-HSD1) (11BETA- HSD1A), [Mouse], complete
AU019271	TC401141	1.265992399 weakly similar to NB1 Glycoprotein (Homo sapiens), partial (85%)
AW547981		1.277883403
AI852776	TC382386	1.278739985 Ku autoantigen (Mus musculus), complete
AW546876	TC375122	1.280703886 Nucleosome assembly protein 1-like protein 4 (Mus musculus), complete
AW545151	TC415940	1.281727527
AA409573	TC424648	1.284631234
AI852578	TC382707	1.287434743 natural resistance associated macrophage protein-2 (Mus musculus), complete
AW556772	TC374744	1.293653077 homologue to N-ACETYLGALACTOSAMINE-6-SULFATASE PRECURSOR (EC 3.1.6.14) (G6S) (GLUCOSAMINE-6-SULFATASE), [Human], partial (95%)
AI853697	TC413774	1.307515385 follistatin-like (Mus musculus), complete
AI852553	TC424648	1.307630929
AW542396	TC408306	1.311321491 TISSUE FACTOR PRECURSOR (TF) (COAGULATION FACTOR III), [Mouse], complete
AI849033	TC372086	1.311971507
AW548558	TC424229	1.313800338
C87282	TC400881	1.324134976 type I transmembrane protein Fn14 (Mus musculus), complete
AI840300	TC368174	1.327711803 Unknown (protein for MGC:13786) (Mus musculus), complete
AU045358	TC390242	1.358492545
AI838985	TC435960	1.370241793 COLD-INDUCIBLE RNA-BINDING PROTEIN (GLYCINE-RICH RNA-BINDING PROTEIN CIRP) (A18 HNRNP), [Mouse, Rat], complete
AU023454		1.38761016
AI851411	TC389898	1.423662281 homologue to von Ebner minor salivary gland protein (Mus musculus), complete
BE380891	TC420954	1.426993114 insulin-like growth factor binding protein 5
AW550895	TC385889	1.445476577 angiotensin II receptor (Mus musculus), complete
AI180560	TC418432	1.459804131 homologue to PERILIPIN (PERI) (LIPID DROPLET-ASSOCIATED PROTEIN), [Rat], partial (50%)
AA240093	TC444003	1.473292844 melusin (Mus musculus), complete
AW545135	TC435817	1.492413102 homologue to D123 (Rattus norvegicus), complete
AW538107	TC378365	1.609216492 cytokeratin (Mus musculus), complete
AW548440		1.615313518
BE532490	TC435848	1.639211745 serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 2 (Mus musculus), complete
AI841876		1.656398622
AI838607	TC374795	1.680832462 THROMBOSPONDIN 1 PRECURSOR, [Mouse], complete
AU040207		1.691398969
AI835238	TC378571	1.691924561 glucose transport protein GT1 - mouse, complete
AW537016	TC407528	1.711236161 CATHEPSIN L PRECURSOR (EC 3.4.22.15) (MAJOR EXCRETED PROTEIN) (MEP), [Mouse], complete
AW551407	TC382731	1.741434023 RIKEN cDNA 2310042P20 gene (Mus musculus), complete
AI847805	TC378384	1.748790824 OSTEOPOINTIN PRECURSOR (BONE SIALOPROTEIN 1) (MINOPONTIN) (EARLY L YMPHOCTYE ACTIVATION 1 PROTEIN) (SECRETED PHOSPHOPROTEIN 1) (SPP-1) (ZAR) (CALCIUM OXALATE CRYSTAL GROWTH INHIBITOR PROTEIN), [Mouse], complete
AI846661	TC378574	1.763197406 putative (Mus musculus), complete
AW538778	TC378465	1.796297477 rbm3 (Mus musculus), complete
AW550637	TC417570	1.801001594 putative (Mus musculus), complete
AW550998		1.821035566
AU041685	TC428469	1.848933106 putative (Mus musculus), complete
AW551313	TC420954	1.852727011 insulin-like growth factor binding protein 5
AI841883		1.87080195
BE361783	TC428235	1.896068608 ORF (Mus musculus), partial (95%)
AW548231	TC392001	1.89970553
AI849214	TC400631	1.917679934 homologue to Similar to transforming growth factor, beta 3 (Mus musculus), partial (33%)
AI322365	TC390516	1.997089386
AI853496	TC418458	2.063736177 Similar to putative heme-binding protein (Mus musculus), complete
AI840126	TC378353	2.07100794
BE377972	TC413729	2.07530521 insulin-like growth factor binding protein-4
AW536807	TC413738	2.078604287 myristylated alanine-rich protein kinase C substrate - mouse, complete
AW551388	TC436345	2.079918369 TRANSCRIPTION FACTOR E2F6 (E2F-6) (E2F-BINDING SITE MODULATING ACTIVITY PROTEIN) (EMA), [Mouse], complete

AW554429	TC404934	2.094645598
AU041376	TC413729	2.0968043 insulin-like growth factor binding protein-4
AW548197	TC401169	2.103156532 calinin beta (Mus musculus), complete
AW549905	TC400269	2.107727642 putative (Mus musculus), complete
AIB45477	TC374923	2.108034996 I-kappa B alpha chain (Mus musculus), complete
AW549905	TC400269	2.11877535 putative (Mus musculus), complete
AW549342	TC422762	2.119464012
AW538349	TC424726	2.124694697 annexin I - mouse, complete
AW546106	TC418146	2.168625188 THYMIDYLATE SYNTHASE (EC 2.1.1.45) (TS), [Mouse], complete
AW549905	TC400269	2.190083165 putative (Mus musculus), complete
AW550525		2.195329398
AU016154	TC413670	2.203799012 CALCICYCLIN (PROLACTIN RECEPTOR ASSOCIATED PROTEIN) (SB10), [Mouse], complete
AW549905	TC400269	2.224160485 putative (Mus musculus), complete
AW551033	TC371950	2.226623651 MEGF8 (Rattus norvegicus), partial (28%)
AW550167		2.228388613
AW550996		2.250539465
AW546817		2.25531098
AW546331	TC408662	2.257616812 homologue to Similar to Myosin of the dilute-myosin-V family (Mus musculus), complete
C79876	TC416505	2.25943572
AW549905	TC400269	2.263126481 putative (Mus musculus), complete
AIB53560	TC424726	2.27744811 annexin I - mouse, complete
AW549190		2.315808526
AW546550		2.356939562
AIB39031		2.374877823
C79775	TC400269	2.376336491 putative (Mus musculus), complete
AIB47573	TC393323	2.376523996 serine protease OMI (Mus musculus), complete
AIB39412	TC378352	2.385857268 ACYL-COA DESATURASE 2 (EC 1.14.99.5) (STEAROYL-COA DESATURASE 2) (FATTY ACID DESATURASE 2) (DELTA(9)-DESATURASE 2), [Mouse], complete
AW548342	TC367590	2.413898893 putative (Mus musculus), complete
C79602	TC378352	2.417041559 ACYL-COA DESATURASE 2 (EC 1.14.99.5) (STEAROYL-COA DESATURASE 2) (FATTY ACID DESATURASE 2) (DELTA(9)-DESATURASE 2), [Mouse], complete
AW549081	TC393225	2.419105637 NUCLEOLYSIN TIA-1 (RNA-BINDING PROTEIN TIA-1), [Mouse], complete
AIB38762	TC393148	2.425604659 putative (Mus musculus), complete
AW539788	TC424943	2.489146629 METALLOTHIONEIN-II (MT-II), [Mouse], complete
AW550072		2.521166332
AW547812		2.542900302
C75991		2.548295101
AW548291	TC393148	2.57796485 putative (Mus musculus), complete
AW536336	TC378352	2.647304867 ACYL-COA DESATURASE 2 (EC 1.14.99.5) (STEAROYL-COA DESATURASE 2) (FATTY ACID DESATURASE 2) (DELTA(9)-DESATURASE 2), [Mouse], complete
AIB44743		2.780386732
AU041878	TC403907	2.859773836 CLUSTERIN PRECURSOR (SULFATED GLYCOPROTEIN 2) (SGP-2) (CLUSTRIN) (APOLIPOPROTEIN J) (APO-J), [Mouse], complete
C86605	TC380202	2.926361582
AU042260	TC425528	3.563305187 COMPLEMENT FACTOR I PRECURSOR (EC 3.4.21.45) (C3B/C4B INACTIVATOR), [Mouse], complete
AIB39302	TC403907	3.910614801 CLUSTERIN PRECURSOR (SULFATED GLYCOPROTEIN 2) (SGP-2) (CLUSTRIN) (APOLIPOPROTEIN J) (APO-J), [Mouse], complete
AIB36624	TC403907	3.970208011 CLUSTERIN PRECURSOR (SULFATED GLYCOPROTEIN 2) (SGP-2) (CLUSTRIN) (APOLIPOPROTEIN J) (APO-J), [Mouse], complete
AU040587		4.213721341

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