

Clone ID	Annotation	E-Value	Log2 Fold Change	KO ID
AOSB412	DNA polymerase epsilon catalytic subunit A	3.53E-23	1.8841	K2324
AOSB472	Ubiquitin carboxyl-terminal hydrolase 5	2.22E-31	1.2676	K11836
AOSB771	Chromodomain-helicase-DNA-binding protein 8	8.75E-07	1.2790	
AOSC409	Predicted protein	2.29E-16	1.7285	
AOSC633	Peptidyl-prolyl cis-trans isomerase FKBP3	6.79E-19	1.7225	
AOSC785	Cytochrome c	1.75E-61	1.7217	K8738
AOSC853	Putative uncharacterized protein	1.58E-64	1.2502	
AOSF463	Ankyrin-1	5.70E-41	1.1217	K16726
AOSF545	C014-D2 G-protein coupled receptor	4.08E-15	1.0784	
AOSF761	Marginal zone B- and B1-cell-specific protein	3.46E-08	1.4082	
AOSF780	Myosin-2 essential light chain	9.09E-25	1.3775	K12751
AOSF817	Predicted protein	7.13E-51	1.4893	
AOSF969	Myosin-VIIa	4.50E-15	1.0064	K1359
AOSF1075	Next to BRCA1 gene 1 protein	7.09E-09	1.2727	
AOSF1088	Putative uncharacterized protein (Fragment)	2.61E-07	0.9375	
AOSF1094	Predicted protein	4.17E-82	0.9219	
AOSF1111	Methionine aminopeptidase 2	9.29E-11	1.7101	K1265
AOSF1131	GFP-like fluorescent chromoprotein cFP484	2.80E-83	0.9599	
AOSF1176	Galaxin	4.72E-35	0.8053	
AOSF1317	Predicted protein	1.01E-10	1.2657	
AOSF1373	Thioredoxin	2.15E-31	1.3360	K3671
AOSF1395	Predicted protein (Fragment)	8.30E-12	1.0989	
AOSF1447	Glutathione S-transferase Mu 5	2.22E-36	1.8475	K799
AOSF1451	Putative heat shock protein HSP 90-alpha A2	9.05E-123	1.4135	K479
AOSF1451	Putative heat shock protein HSP 90-alpha A2	2.56E-122	1.4135	K479
AOSF1460	Probable E3 ubiquitin-protein ligase HERC1	1.65E-31	1.0439	K1594
AOSF1482	Fanconi anemia, complementation group D2 (Fragment)	2.61E-09	1.1184	
AOSF1484	S-adenosylmethionine synthase	1.90E-37	1.6865	K789
AOSF1513	Uncharacterized protein	1.64E-15	1.6517	
AOSF1525	Uncharacterized protein	3.34E-10	1.2263	
CAGI2339	Uncharacterized protein	4.66E-29	1.4150	
CAGI2374	Mitogen-activated protein kinase 15	4.04E-55	1.4286	K8293
CAGI2441	CCAAT/enhancer-binding protein gamma	3.46E-19	1.1086	
CAGI2444	Dynein gamma chain, flagellar outer arm	3.50E-36	0.9206	
CAGI2506	Putative uncharacterized protein	4.79E-06	2.8634	
CAGI2542	Putative uncharacterized protein (Fragment)	1.20E-06	1.0813	
CAGI2742	Uncharacterized protein	2.57E-18	1.2916	
CAGI2763	Putative uncharacterized protein	3.95E-21	0.8680	
CAGI2773	Fibropellin-1	4.08E-19	2.2161	
CAGI2784	Putative uncharacterized protein	5.10E-14	1.9599	
CAGI2787	Putative uncharacterized protein	8.03E-12	3.0074	
CAGI2801	Putative uncharacterized protein	1.06E-25	1.3495	
CAGI2815	Putative uncharacterized protein	5.09E-44	2.3032	
CAGI2833	Putative uncharacterized protein	1.60E-32	1.0146	
CAGI2842	Putative uncharacterized protein (Fragment)	3.94E-10	2.7325	
CAGI2859	Putative uncharacterized protein	9.36E-42	1.9755	
CAGI2868	Putative uncharacterized protein	4.59E-08	2.1964	
CAGI2884	Uncharacterized protein	4.42E-18	1.1515	
CAGI2964	Retinal guanylyl cyclase 2	9.59E-16	1.9749	K12321
CAGI2990	Putative uncharacterized protein	6.57E-32	0.9876	
CAGI2991	Uncharacterized protein	2.64E-16	1.9545	
CAGI3009	Halomucin	2.95E-06	1.8127	
CAGI3016	Putative uncharacterized protein	5.32E-09	1.6888	
CAGI3050	Putative uncharacterized protein	1.28E-27	1.4744	
CAON422	Neutral alpha-glucosidase AB	1.83E-32	1.9069	K5546

CAON453	Kinesin-like protein unc-104	1.05E-39	1.0018	K17914
CAON500	Ubiquitin carboxyl-terminal hydrolase 24	3.87E-28	0.9610	K1184
CAON612	Activating transcription factor 4-like protein	2.09E-07	1.6989	
CAON1102	Iron/zinc purple acid phosphatase-like protein	1.73E-06	1.0219	
CAON1209	60S ribosomal protein L44 Q	2.19E-54	1.0928	K2929
CAON1226	LWamide neuropeptides	5.08E-07	1.0148	
CAON1304	Uncharacterized protein	2.03E-07	1.0197	
CAON1314	DNA topoisomerase I, mitochondrial	2.59E-44	1.1344	K3163
CAON1404	Alanine aminotransferase 2	4.15E-29	2.5369	K814
CAON1485	Predicted protein	3.72E-10	1.1345	
CAON1582	Solute carrier organic anion transporter family member 4A1	5.96E-45	1.5169	K14354
CAON1646	Nucleoside diphosphate kinase B	9.75E-78	1.5853	K94
CAON1781	Transposon TX1 uncharacterized 149 kDa protein	5.36E-10	1.5032	
CAON1787	Putative uncharacterized protein (Fragment)	7.28E-06	1.2428	
CAOO902	Elongation factor 1-alpha 2	1.41E-23	1.1064	
CAOO1946	Fanconi anemia group I protein homolog	4.99E-48	0.9906	
CAOO2018	Heat shock protein 105 kDa	4.08E-11	0.8565	K9485
CAOO2241	Histone acetyltransferase Tip60	2.64E-44	0.9124	K1134
CAOO2584	Mitogen-activated protein kinase 1	1.33E-71	0.9664	K4371
CAOO2593	Ras-related protein SEC4	4.91E-13	1.1521	
CCHW786	Lipoma HMGIC fusion partner homolog	1.24E-46	1.7755	
CCHW795	Uncharacterized protein	2.27E-09	1.2961	
CCHW841	Transmembrane emp24 domain-containing protein 4	5.17E-101	0.9655	
CCHW842	Predicted protein	1.43E-32	0.8725	
CCHW854	WD repeat-containing protein 66	7.91E-62	1.1238	
CCHW875	Predicted protein (Fragment)	9.42E-46	1.3319	
CCHW936	Mediator of RNA polymerase II transcription subunit 4	8.70E-33	1.2615	K15146
CCHW1049	Serine hydrolase-like protein	1.34E-36	2.0138	
CCHW1070	Trans-Golgi network integral membrane protein 2	1.42E-07	1.2688	
CCHW1137	Uncharacterized protein	4.59E-07	0.9559	
CCHW1161	Zinc finger protein 36, C3H1 type-like 1	2.17E-43	0.8913	
CCHW1163	Predicted protein	4.02E-30	1.1884	
CCHW1174	Protein salvador homolog 1	1.07E-36	1.3112	K16686
CCHW1181	Sushi, von Willebrand factor type A, EGF and pentraxin domain-containing protein 1	1.17E-19	1.4415	K16681
CCHW1182	Protein bicaudal C homolog 1-A	1.04E-20	1.4520	
CCHW1183	40S ribosomal protein S15	1.84E-63	0.8106	K2958
CCHW1190	Spermatogenesis-defective protein 39 homolog	7.14E-49	1.0124	
CCHW1207	Coiled-coil domain-containing protein 166	4.36E-09	1.8303	
CCHW1209	Forkhead box protein O1-B	1.11E-17	1.5077	K721
CCHW1217	Multiple coagulation factor deficiency protein 2 homolog	1.14E-28	1.0281	
CCHW1225	Centrosomal protein of 78 kDa	8.77E-06	1.4335	
CCHW1232	Putative uncharacterized protein	1.89E-24	1.3593	
CCHW1233	Steroid 17-alpha-hydroxylase/17,20 lyase	4.40E-19	1.6079	K14985
CCHW1239	Predicted protein	1.65E-13	2.0579	
CCHW1255	Mitochondrial import inner membrane translocase subunit Tim8 A	3.44E-19	1.0426	K1778
CCHW1259	Dihydrolipoyllysine-residue succinyltransferase component of 2-oxoglutarate dehydrogenase complex, mitochondrial	1.62E-62	1.5837	K658
CCHW1277	Uncharacterized protein	5.85E-14	0.9682	
CCHW1279	Mitochondrial ornithine transporter 1	5.42E-08	1.0369	
CCHW1285	Predicted protein (Fragment)	2.03E-84	1.4985	
CCHW1288	Thioredoxin, mitochondrial	1.24E-27	1.8210	K3671
CCHW1303	Akirin-2	1.66E-29	1.1140	
CCHW1307	Histidine triad nucleotide-binding protein 1	1.64E-51	1.1936	K253
CCHW1316	Tripartite motif-containing protein 3	3.11E-17	1.8416	K1235
CCHW1324	Guanine nucleotide-binding protein G(q) subunit alpha	1.69E-52	1.3573	K4634
CCHW1335	Predicted protein	3.34E-32	0.9885	

CCHW1351	Predicted protein (Fragment)	4.39E-25	1.1460	
CCHW1367	Tetratricopeptide repeat protein 9C	3.42E-25	1.0607	
CCHW1371	Nuclear distribution protein nudE-like 1	3.96E-55	2.3615	K16738
CCHW1386	Protein disulfide-isomerase 1	1.03E-08	0.9048	
CCHW1393	Growth/differentiation factor 8	9.06E-11	1.1560	K5497
CCHW1401	Ras-related and estrogen-regulated growth inhibitor	1.29E-17	1.3056	K7855
CCHW1410	WAP four-disulfide core domain protein 18 (Fragment)	7.53E-08	1.6576	
CCHW1463	Inositol hexakisphosphate kinase 1	3.96E-50	1.4501	K7756
CCHW1467	Cytochrome c oxidase subunit 5A, mitochondrial	1.46E-44	0.9871	K2264
CCHW1581	Putative L-aspartate dehydrogenase	1.55E-41	0.9729	
CCHW1596	Predicted protein	1.85E-09	1.2703	
CCHW1609	Predicted protein	1.15E-09	1.1060	
CCHW1614	Predicted protein	1.10E-45	0.8661	
CCHW1632	Serine--pyruvate aminotransferase, mitochondrial	3.10E-68	1.6829	K83
CCHW1636	Putative rte all (Fragment)	7.37E-11	0.9106	
CCHW1684	Probable C-mannosyltransferase DPY19L3	5.64E-12	1.5704	
CCHW1723	Transcription factor HES-1	1.12E-13	0.9855	K654
CCHW1775	Uncharacterized protein	7.78E-12	1.9582	
CCHW1796	Golgi apparatus protein 1	1.22E-24	0.9386	K6816
CCHW1822	Uncharacterized protein	6.81E-23	1.2675	
CCHW1823	Transmembrane emp24 domain-containing protein 1	2.74E-27	1.2737	
CCHW1826	Signal recognition particle 54 kDa protein	9.96E-19	1.4398	K316
CCHW1849	Egg protein	5.26E-47	0.8663	
CCHW1871	Gamma-soluble NSF attachment protein	3.68E-53	1.0702	
CCHW1878	NADH dehydrogenase [ubiquinone] iron-sulfur protein 5-A	7.39E-09	0.8423	
CCHW1889	Arylsulfatase B	5.17E-36	2.5649	K1135
CCHW1915	Predicted protein	3.14E-37	1.7147	
CCHW1922	Uncharacterized protein	7.52E-48	0.8975	
CCHW1947	Histone H2B 1/2	1.39E-58	1.2594	K11252
CCHW1950	Thioredoxin reductase 3	2.10E-62	1.3673	K3676
CCHW1958	A disintegrin and metalloproteinase with thrombospondin motifs 17	3.45E-07	0.8884	
CCHW1986	Plasma kallikrein	2.16E-43	1.0409	
CCHW1986	Plasminogen (Fragment)	1.87E-27	1.0409	
CCHW2000	Predicted protein	2.62E-25	0.9707	
CCHW2013	COG: Uncharacterized conserved protein	1.96E-08	1.1904	
CCHW2042	Putative uncharacterized protein	5.21E-43	1.2848	
CCHW2072	Protein 4.1	7.44E-69	1.1908	K617
CCHW2074	Chromosome undetermined scaffold_105, whole genome shotgun sequence	1.99E-42	0.8232	
CCHW2116	Cdc42 homolog	3.07E-47	0.9055	K4393
CCHW2130	Predicted protein (Fragment)	5.42E-15	1.4329	
CCHW2133	Proteasome inhibitor PI31 subunit	1.38E-09	1.2259	
CCHW2139	D-amino-acid oxidase	5.92E-08	1.2970	
CCHW2142	Membrane progesterin receptor gamma-A	7.05E-19	0.9183	
CCHW2168	Predicted protein	5.43E-19	1.7220	
CCHW2186	Thymosin beta-15B	1.39E-07	1.2623	
CCHW2187	Predicted protein	2.98E-40	1.2161	
CCHW2193	Zinc finger protein 330 homolog	4.28E-62	1.2388	
CCHW2208	Predicted protein	1.23E-12	0.8648	
CCHW2215	Myosin regulatory light polypeptide 9	1.00E-61	1.3390	K12757
CCHW2274	Nucleoside diphosphate kinase B	1.18E-81	0.9897	K94
CCHW2296	Sodium-dependent phosphate transport protein 2B	8.46E-48	1.7188	K14683
CCHW2328	RNA-binding protein 1	2.91E-21	1.2177	K12896
CCHW2348	Chymotrypsinogen B	6.21E-11	1.1099	
CCHW2374	Ragulator complex protein LAMTOR4 homolog	5.63E-21	0.7871	
CCHW2377	U3 small nucleolar RNA-interacting protein 2	5.65E-64	1.2610	K14793
CCHW2382	ATP-binding cassette sub-family E member 1	1.97E-30	1.4962	K6174

CCHW2403	Elongation factor 1-alpha	1.18E-120	1.3253	K3231
CCHW2406	Proteasome assembly chaperone 1	3.17E-12	1.3564	
CCHW2424	Hsc70-interacting protein	2.43E-34	1.3281	K956
CCHW2427	Predicted protein	1.09E-16	1.4710	
CCHW2459	46 kDa FK506-binding nuclear protein	6.45E-18	2.0149	
CCHW2485	Renin receptor	2.40E-08	1.0846	
CCHW2496	Parkin coregulated gene protein homolog	9.56E-87	1.4311	
CCHW2500	Protein argonaute-2	7.77E-109	1.2474	K11593
CCHW2513	U1 small nuclear ribonucleoprotein 70 kDa	1.25E-43	1.1785	K1193
CCHW2529	Charged multivesicular body protein 3	1.90E-26	1.0733	K12193
CCHW2561	Phosphatase and actin regulator 1	4.72E-11	0.9631	
CCHW2572	Tubulin alpha chain	1.13E-145	1.1496	K7374
CCHW2602	Methionine-R-sulfoxide reductase B1	1.97E-34	1.2540	K735
CCHW2655	Prefoldin subunit 6	3.19E-27	1.7249	K4798
CCHW2661	26S proteasome non-ATPase regulatory subunit 7	2.69E-82	1.0256	K338
CCHW2762	Predicted protein	4.96E-17	1.5310	
CCHW2830	Melatonin receptor type 1B-B	9.56E-06	1.0760	
CCHW2857	Pancreas transcription factor 1 subunit alpha	1.14E-13	1.1916	K973
CCHW2880	E3 ubiquitin-protein ligase HECW2	3.98E-34	1.1713	K12168
CCHW2914	E3 ubiquitin-protein ligase RING2-A	2.79E-92	0.9810	K1695
CCHW3027	Predicted protein	8.36E-14	1.5948	
CCHW3036	P-type proton ATPase	1.12E-98	1.0440	
CCHW3060	Phenylalanine--tRNA ligase beta subunit	5.51E-76	1.3143	K189
CCHW3088	COP9 signalosome complex subunit 5	4.25E-123	1.8065	K9613
CCHW3097	Peptidase M20 domain-containing protein 2	6.89E-09	1.7005	
CCHW3115	Flavin reductase (NADPH)	3.12E-22	1.4172	K591
CCHW3125	Predicted protein (Fragment)	4.44E-45	1.1351	
CCHW3127	33 kDa inner dynein arm light chain, axonemal	5.04E-24	1.9144	K141
CCHW3130	Lipase member I	9.97E-48	0.8546	K1475
CCHW3260	Pro-epidermal growth factor	6.28E-22	1.4091	K6825
CCHW3282	TM2 domain-containing protein 1	1.93E-49	1.0504	
CCHW3309	Cleavage and polyadenylation specificity factor subunit 6	5.29E-31	1.5876	K14398
CCHW3334	60S ribosomal protein L10a	8.93E-112	1.1497	K2865
CCHW3373	Spliceosome RNA helicase Ddx39b	1.11E-114	1.3642	K12812
CCHW3377	Hemagglutinin/amebocyte aggregation factor	5.73E-12	1.1901	
CCHW3464	Radixin	1.04E-82	1.0694	K5762
CCHW3584	STE20-like serine/threonine-protein kinase	1.39E-63	1.2297	K8836
CCHW3870	Ras-related protein Rab-21	2.13E-24	1.1153	
CCHW4042	Scleractinian cysteine-rich protein	1.21E-32	0.9483	
CCHW4129	Cystatin-A1	8.05E-25	1.2242	K1397
CCHW4181	Predicted protein	1.09E-26	0.8819	
CCHW4218	Glutaryl-CoA dehydrogenase, mitochondrial	2.27E-103	1.3381	K252
CCHW4272	10 kDa heat shock protein, mitochondrial	1.37E-36	1.2152	K478
CCHW4347	Signal peptidase complex subunit 3	1.14E-64	1.0557	K12948
CCHW4490	D-tyrosyl-tRNA(Tyr) deacylase 1	2.54E-62	1.0023	K756
CCHW4512	G patch domain-containing protein 4	6.61E-47	1.4971	
CCHW4528	Clathrin heavy chain 1	9.11E-144	0.9299	K4646
CCHW4636	Heat shock factor-binding protein 1	7.23E-10	1.4687	
CCHW4711	Tenascin-R	6.16E-60	1.3857	K6252
CCHW4732	Ubiquitin-conjugating enzyme E2 Z	6.86E-104	1.0467	K1585
CCHW4770	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransferase	3.02E-08	1.9446	
CCHW4833	72 kDa type IV collagenase	3.38E-22	1.1214	K82
CCHW4873	Radial spoke head protein 6 homolog A	5.52E-57	0.9459	
CCHW5049	F-actin-capping protein subunit alpha	6.16E-97	1.2080	K1364
CCHW5057	Universal stress protein A-like protein	4.23E-12	1.9465	
CCHW5186	28S ribosomal protein S21, mitochondrial	1.33E-18	1.2937	K297
CCHW5228	Rhamnose-binding lectin	1.24E-21	1.0931	K4592

CCHW5393	Reactive oxygen species modulator 1	9.88E-26	1.6827	
CCHW5749	Ankyrin repeat and IBR domain-containing protein 1	5.09E-80	2.1550	K11967
CCHW5764	U6 snRNA-associated Sm-like protein LSm1	1.02E-45	1.1668	K1262
CCHW6093	IQ and ubiquitin-like domain-containing protein	1.96E-43	1.4465	
CCHW6576	Heat shock cognate 71 kDa protein	4.74E-22	1.1409	K3283
CCHW6627	Mothers against decapentaplegic homolog 3	6.48E-54	1.2837	K45
CCHW6699	Uncharacterized protein	1.28E-06	1.3211	
CCHW6705	BolA-like protein 2	4.85E-22	0.7384	
CCHW6709	Predicted protein	9.36E-27	1.1261	
CCHW6764	Predicted protein	7.55E-30	1.2382	
CCHW6782	DNA-directed RNA polymerases I, II, and III subunit RPABC4	6.90E-21	1.8426	K39
CCHW6812	Ribonucleoside-diphosphate reductase subunit M2	1.08E-82	1.2212	K188
CCHW6864	Group XIIB secretory phospholipase A2-like protein	1.13E-08	1.1278	
CCHW6865	Putative ankyrin repeat protein FPV219	1.59E-09	1.1372	
CCHW6903	Sugar transporter ERD6-like 4	8.78E-36	0.8029	K8145
CCHW6936	Splicing factor 3A subunit 2	8.75E-117	1.2878	K12826
CCHW7314	Proteasome activator complex subunit 4B	2.03E-09	1.1243	
CCHW7412	Neurexin-4	1.04E-34	1.6895	
CCHW7562	Thioredoxin	1.33E-37	0.9969	K3671
CCHW7667	V-type proton ATPase subunit G	2.94E-36	0.9753	K2152
CCHW7818	Acyl-CoA dehydrogenase family member 10	2.44E-11	0.8324	
CCHW7988	Probable cation-transporting ATPase 13A1	7.58E-30	1.9100	K1495
CCHW7992	Exportin-6	6.06E-46	1.0032	
CCHW8046	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 5	6.76E-38	0.9803	K3949
CCHW8127	Ras-related protein Rab-3	1.02E-132	0.9543	K7883
CCHW8167	DNA-directed RNA polymerase III subunit RPC9	1.62E-29	0.9294	
CCHW8171	Peripheral-type benzodiazepine receptor-associated protein 1	3.19E-16	0.9923	K17591
CCHW8173	Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial	7.12E-56	1.2081	K12663
CCHW8189	39S ribosomal protein L34, mitochondrial	5.34E-13	0.9619	
CCHW8200	Glutathionyl-hydroquinone reductase YqjG	5.78E-55	1.8857	K7393
CCHW8224	Putative uncharacterized protein	3.86E-17	0.8415	
CCHW8248	Selenoprotein	2.24E-21	0.9730	
CCHW8251	Peptidyl-prolyl cis-trans isomerase FKBP1A	1.09E-32	0.9563	K9568
CCHW8283	Pre-mRNA-processing factor 6	1.41E-97	1.1580	K12855
CCHW8288	Breast cancer anti-estrogen resistance protein 3 homolog	5.24E-10	1.7510	
CCHW8301	Golgi-associated plant pathogenesis-related protein 1	2.62E-20	1.2480	
CCHW8343	Predicted protein	2.23E-13	0.9432	
CCHW8403	Platelet-activating factor acetylhydrolase IB subunit beta	7.18E-11	1.4413	
CCHW8409	Mitotic spindle-associated MMXD complex subunit MIP18	4.38E-77	1.5699	
CCHW8414	Nuclear pore complex protein Nup205	1.27E-23	1.3792	K1431
CCHW8464	Tudor domain-containing protein 1	4.21E-13	1.9719	K1845
CCHW8469	Protein fem-1 homolog A	1.00E-08	1.2466	
CCHW8656	Nucleolar protein 6	7.05E-54	1.2969	K14544
CCHW8675	Tetratricopeptide repeat protein 40	4.31E-12	0.9088	
CCHW8751	Dynactin subunit 6	3.74E-97	1.4514	K1428
CCHW8763	X-ray radiation resistance-associated protein 1	5.90E-06	1.2223	
CCHW8816	Peroxisomal membrane protein PMP34	1.23E-18	1.2087	K13354
CCHW8920	Ragulator complex protein LAMTOR5 homolog	2.93E-40	0.8049	K16344
CCHW9612	Uncharacterized protein	1.59E-14	0.8412	
CCHW9616	1-acylglycerol-3-phosphate O=-acyltransferase ABHD5	6.35E-79	1.2657	
CCHW9622	Double-stranded RNA-binding protein 2	9.71E-09	1.7434	
CCHW9821	Cytochrome c oxidase subunit 6B1	6.26E-26	1.3848	K2267
CCHW9841	Tetratricopeptide repeat protein 26	3.05E-152	1.2238	
CCHW9873	Uracil phosphoribosyltransferase homolog	2.66E-29	1.5982	K761
CCHW9886	Tyrosine-protein kinase isoform SRK4	4.15E-07	1.1703	
CCHW9927	GFP-like fluorescent chromoprotein cFP484	8.12E-30	1.2423	
CCHW9980	Predicted protein	4.63E-70	0.9198	

CCHW10396	Coiled-coil domain-containing protein lobo homolog	1.04E-53	1.0134	
CCHW10464	Dihydropyrimidinase-related protein 2	4.07E-09	1.4788	
CCHW10491	Predicted protein	4.15E-11	1.4284	
CCHW10707	Predicted protein	7.10E-46	1.3768	
CCHW10745	1-acylglycerol-3-phosphate O ⁼ -acyltransferase ABHD5	1.25E-88	0.9013	
CCHW11014	Glutamyl aminopeptidase	7.54E-30	1.1620	K11141
CCHW11287	C-Myc-binding protein	1.16E-30	0.8252	
CCHW11644	F-box/LRR-repeat protein 13	5.73E-21	1.5343	K1268
CCHW11647	PCTP-like protein	4.68E-35	1.5475	
CCHW11698	Homeobox protein OTX2-B	3.80E-29	1.4888	K9326
CCHW11702	Peroxidasin	6.78E-07	1.4279	
CCHW11709	Intraflagellar transport protein 27 homolog	8.48E-74	1.2985	K7934
CCHW11728	Cysteine/serine-rich nuclear protein 3	1.15E-27	1.4394	K17494
CCHW11836	Fibulin-1	1.55E-19	0.9516	K1737
CCHW11870	Egl nine homolog 1	6.04E-36	0.9604	K9592
CCHW12017	Predicted protein	5.26E-13	0.9769	
CCHW12026	Splicing factor 3A subunit 3	1.04E-83	0.9723	K12827
CCHW12150	EF-hand calcium-binding domain-containing protein 6 (Fragment)	2.08E-23	1.0838	
CCHW12230	Transmembrane protein 241	3.05E-37	0.9769	
CCHW12237	Peptidyl-prolyl cis-trans isomerase FKBP2	1.18E-61	1.1054	K9569
CCHW12679	Ribosomal RNA-processing protein 7 homolog A	2.28E-45	1.4265	K14545
CCHW12680	Phytanoyl-CoA dioxygenase, peroxisomal	4.93E-15	1.0413	K477
CCHW12682	Nuclear export mediator factor NEMF	1.00E-31	2.1299	
CCHW12687	Transcription initiation factor TFIID subunit 13	1.66E-50	1.3056	K3127
CCHW12694	Zinc metalloproteinase nas-14	8.87E-46	1.4077	K876
CCHW12729	Protease inhibitor LmKTT-1a	1.46E-06	1.5204	
CCHW12732	Anionic trypsin	1.15E-49	1.2278	K1324
CCHW12754	Putative uncharacterized protein (Fragment)	2.24E-06	1.8596	
CCHW12760	Calcium/calmodulin-dependent protein kinase type 1D	4.36E-110	1.0756	K8794
CCHW12771	Uncharacterized protein (Fragment)	2.59E-23	1.3023	
CCHW12787	Putative uncharacterized protein	4.60E-43	0.9209	
CCHW12794	Transmembrane protein 256 homolog	2.00E-20	1.7855	
CCHW12802	Protein FAM76B	8.77E-49	1.2496	
CCHW12803	Predicted protein	8.55E-40	1.4654	
CCHW12826	Cathepsin B	1.66E-29	1.0405	K1363
CCHW12842	15-hydroxyprostaglandin dehydrogenase [NAD(+)]	1.46E-56	1.1671	K59
CCHW12847	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1	3.04E-102	0.8555	K4536
CCHW12873	Achaete-scute homolog 4	1.71E-16	1.4825	K967
CCHW12970	40S ribosomal protein S2-3	5.07E-81	1.8404	
CCHW12976	Predicted protein	9.86E-72	1.4318	K13755
CCHW12988	Thiosulfate sulfurtransferase/rhodanese-like domain-containing protein 3	1.89E-25	1.3459	
CCHW12990	Thyrotroph embryonic factor	5.12E-09	0.9278	
CCHW13029	Solute carrier family 22 member 18	1.51E-31	1.2054	K8214
CCHW13060	Abhydrolase domain-containing protein 4	2.65E-25	2.0208	
CCHW13115	Diphosphoinositol polyphosphate phosphohydrolase 1	2.23E-39	1.0206	K7766
CCHW13248	Predicted protein	1.12E-12	1.1730	
CCHW13432	STE20/SPS1-related proline-alanine-rich protein kinase	5.49E-108	0.8427	K8835
CCHW16981	Ancient ubiquitous protein 1	6.70E-21	1.0417	
CCHW17067	Protein NLRC5	3.16E-07	1.1076	
CCHW17448	Protein C-ets-2 (Fragment)	1.39E-32	0.9091	K2678