

Table S7.1. Transcript analysis of adult male vs adult female gene expression.

GeneID	Gene_length	Male reads	Female reads	Male-RPKM	Female-RPKM	log2 Ratio(F/M)	Up-Down-Regulation(F/M)	P-value	FDR	Description
CCG008720.1	324	10	0	1.32622638	0.001	-10.37311134	Down	0.0002961	0.00047146	AAEL006662-PA OS=Aedes aegypti GN=AAEL006662 PE=4 SV=1
CCG001977.1	372	10	0	1.15510039	0.001	-10.17380253	Down	0.0002961	0.00047158	AAEL007682-PA OS=Aedes aegypti GN=AAEL007682 PE=4 SV=1
CCG001443.1	372	10	0	1.15510039	0.001	-10.17380253	Down	0.0002961	0.00047155	AAEL003256-PA OS=Aedes aegypti GN=AAEL003256 PE=4 SV=1
CCG008172.1	495	10	0	0.86807545	0.001	-9.761676626	Down	0.0002961	0.00047152	-
CCG018587.1	513	10	0	0.83761666	0.001	-9.710146326	Down	0.0002961	0.0004714	AAEL005966-PA OS=Aedes aegypti GN=AAEL005966 PE=4 SV=1
CCG016364.1	552	10	0	0.77843722	0.001	-9.604436884	Down	0.0002961	0.00047143	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003311 PE=4 SV=1
CCG024454.1	617	10	0	0.69643006	0.001	-9.443834662	Down	0.0002961	0.00047167	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC006955 PE=4 SV=1
CCG006445.1	720	10	0	0.59680187	0.001	-9.221108245	Down	0.0002961	0.00047125	AAEL009771-PA OS=Aedes aegypti GN=AAEL009771 PE=3 SV=1
CCG019346.1	726	10	0	0.59186962	0.001	-9.209135603	Down	0.0002961	0.00047161	AAEL001256-PA OS=Aedes aegypti GN=AAEL001256 PE=4 SV=1
CCG026240.1	762	10	0	0.56390728	0.001	-9.139314154	Down	0.0002961	0.00047128	AAEL009564-PA OS=Aedes aegypti GN=AAEL009564 PE=4 SV=1
CCG024932.1	789	10	0	0.54461007	0.001	-9.089079851	Down	0.0002961	0.00047149	AAEL011450-PA OS=Aedes aegypti GN=AAEL011450 PE=3 SV=1
CCG001967.1	1140	10	0	0.3769275	0.001	-8.558143232	Down	0.0002961	0.00047164	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001348 PE=4 SV=1
CCG009416.1	1200	10	0	0.35808112	0.001	-8.484142651	Down	0.0002961	0.00047131	AAEL002916-PA OS=Aedes aegypti GN=AAEL002916 PE=4 SV=1

CCG018326.1	1785	10	0	0.2407268	0.001	-7.911252982	Down	0.0002961	0.00047134	AAEL003976-PA OS=Reedes aegypti GN=AAEL003976 PE=4 SV=1
CCG013652.1	4243	10	0	0.10127206	0.001	-6.662092378	Down	0.0002961	0.00047137	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC012952 PE=4 SV=1
CCG011264.1	345	11	0	1.37004951	0.001	-10.42001231	Down	0.0001328	0.00021562	AAEL008251-PA OS=Reedes aegypti GN=AAEL008251 PE=4 SV=1
CCG005571.1	417	11	0	1.1334942	0.001	-10.14656129	Down	0.0001328	0.00021557	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009105 PE=4 SV=1
CCG021062.1	459	11	0	1.02977577	0.001	-10.00811452	Down	0.0001328	0.00021559	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010139 PE=4 SV=1
CCG021466.1	488	11	0	0.96858008	0.001	-9.919727527	Down	0.0001328	0.00021564	-
CCG015216.1	534	11	0	0.88514435	0.001	-9.789768933	Down	0.0001328	0.00021565	AAEL000085-PA OS=Reedes aegypti GN=AAEL000085 PE=4 SV=1
CCG008792.1	568	11	0	0.83216035	0.001	-9.700717745	Down	0.0001328	0.00021566	-
CCG016091.1	780	11	0	0.60598344	0.001	-9.243134551	Down	0.0001328	0.00021561	AAEL003311-PA OS=Reedes aegypti GN=AAEL003311 PE=4 SV=1
CCG009792.1	1857	11	0	0.25453262	0.001	-7.991706765	Down	0.0001328	0.00021558	AGAP013374-PA OS=Anopheles gambiae GN=AgaP_AGAP013374 PE=4 SV=1
CCG017791.1	288	12	0	1.79040561	0.001	-10.80607075	Down	5.96E-05	9.92E-05	AAEL006722-PA OS=Reedes aegypti GN=AAEL006722 PE=4 SV=1
CCG027620.1	420	12	0	1.2277067	0.001	-10.26175023	Down	5.96E-05	9.91E-05	AAEL008967-PA OS=Reedes aegypti GN=AAEL008967 PE=4 SV=1
CCG022164.1	450	12	0	1.14585959	0.001	-10.16221456	Down	5.96E-05	9.92E-05	AAEL013722-PA OS=Reedes aegypti GN=AAEL013722 PE=4 SV=1
CCG021342.1	552	12	0	0.93412467	0.001	-9.86747129	Down	5.96E-05	9.92E-05	AAEL012690-PA OS=Reedes aegypti GN=AAEL012690 PE=4 SV=1
CCG011393.1	570	12	0	0.90462599	0.001	-9.821177638	Down	5.96E-05	9.92E-05	AAEL000645-PA OS=Reedes aegypti GN=AAEL000645 PE=4 SV=1

CCG006007.1	755	12	0	0.68296267	0.001	-9.415662913	Down	5.96E-05	9.92E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005028 PE=4 SV=1
CCG008062.1	767	12	0	0.67227746	0.001	-9.39291298	Down	5.96E-05	9.91E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018868 PE=4 SV=1
CCG018582.1	819	12	0	0.62959318	0.001	-9.298276105	Down	5.96E-05	9.92E-05	AAEL001330-PA OS=Aedes aegypti GN=AAEL001330 PE=3 SV=1
CCG017551.1	855	12	0	0.60308399	0.001	-9.236215137	Down	5.96E-05	9.92E-05	AAEL009488-PA OS=Aedes aegypti GN=AAEL009488 PE=4 SV=1
CCG000905.1	856	12	0	0.60237946	0.001	-9.234528761	Down	5.96E-05	9.92E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG025374.1	1015	12	0	0.50801657	0.001	-8.988731735	Down	5.96E-05	9.92E-05	-
CCG025421.1	1539	12	0	0.33504666	0.001	-8.388218231	Down	5.96E-05	9.92E-05	AAEL004647-PA OS=Aedes aegypti GN=AAEL004647 PE=4 SV=1
CCG015653.1	1725	12	0	0.29891989	0.001	-8.223615101	Down	5.96E-05	9.92E-05	AAEL014010-PA (Fragment) OS=Aedes aegypti GN=AAEL014010 PE=4 SV=1
CCG024944.1	1749	12	0	0.29481808	0.001	-8.203681173	Down	5.96E-05	9.92E-05	AAEL009646-PA OS=Aedes aegypti GN=AAEL009646 PE=4 SV=1
CCG004710.1	348	13	0	1.60519124	0.001	-10.64852947	Down	2.67E-05	4.57E-05	AAEL009000-PA OS=Aedes aegypti GN=AAEL009000 PE=4 SV=1
CCG007130.1	375	13	0	1.48961747	0.001	-10.54072618	Down	2.67E-05	4.57E-05	AAEL005810-PA OS=Aedes aegypti GN=AAEL005810 PE=4 SV=1
CCG014551.1	445	13	0	1.25529562	0.001	-10.29381144	Down	2.67E-05	4.57E-05	-
CCG021485.1	630	13	0	0.88667706	0.001	-9.792264946	Down	2.67E-05	4.57E-05	-
CCG028100.1	657	13	0	0.85023828	0.001	-9.731723404	Down	2.67E-05	4.57E-05	AAEL008168-PA OS=Aedes aegypti GN=AAEL008168 PE=4 SV=1
CCG002844.1	387	14	0	1.55446068	0.001	-10.60219841	Down	1.20E-05	2.09E-05	-
CCG008953.1	663	14	0	0.90735488	0.001	-9.825523108	Down	1.20E-05	2.09E-05	AAEL010201-PA OS=Aedes aegypti GN=AAEL010201 PE=4 SV=1
CCG004917.1	723	14	0	0.83205572	0.001	-9.700536331	Down	1.20E-05	2.09E-05	-
CCG016788.1	753	14	0	0.79890609	0.001	-9.641882114	Down	1.20E-05	2.09E-05	AAEL002234-PA OS=Aedes aegypti GN=AAEL002234 PE=4 SV=1

CCG020823.1	393	15	0	1.64006621	0.001	-10.67953834	Down	5.38E-06	9.55E-06	-
CCG008356.1	438	15	0	1.47156625	0.001	-10.52313678	Down	5.38E-06	9.55E-06	AAEL012061-PA OS=Aedes aegypti GN=AAEL012061 PE=4 SV=1
CCG013328.1	459	15	0	1.40423969	0.001	-10.4555735	Down	5.38E-06	9.55E-06	AAEL005034-PA OS=Aedes aegypti GN=AAEL005034 PE=3 SV=1
CCG021896.1	369	16	0	1.86318632	0.001	-10.86355624	Down	2.41E-06	4.35E-06	AAEL003177-PA OS=Aedes aegypti GN=AAEL003177 PE=4 SV=1
CCG006970.1	384	16	0	1.79040561	0.001	-10.80607075	Down	2.41E-06	4.35E-06	AAEL010657-PA OS=Aedes aegypti GN=AAEL010657 PE=4 SV=1
CCG012027.1	414	16	0	1.66066607	0.001	-10.69754629	Down	2.41E-06	4.35E-06	AAEL013513-PA OS=Aedes aegypti GN=AAEL013513 PE=4 SV=1
CCG008326.1	1002	16	0	0.68614347	0.001	-9.422366453	Down	2.41E-06	4.35E-06	AAEL009098-PA OS=Aedes aegypti GN=AAEL009098 PE=4 SV=1
CCG027791.1	2109	16	0	0.32599135	0.001	-8.348689867	Down	2.41E-06	4.35E-06	AAEL003321-PA OS=Aedes aegypti GN=AAEL003321 PE=4 SV=1
CCG008230.1	474	17	0	1.54110863	0.001	-10.58975284	Down	1.08E-06	1.99E-06	-
CCG001665.1	1071	17	0	0.68205928	0.001	-9.413753323	Down	1.08E-06	1.99E-06	AAEL002911-PA (Fragment) OS=Aedes aegypti GN=AAEL002911 PE=4 SV=1
CCG009531.1	1125	17	0	0.64932043	0.001	-9.342786801	Down	1.08E-06	1.99E-06	AAEL006484-PA OS=Aedes aegypti GN=GPROP3 PE=3 SV=1
CCG027452.1	1170	17	0	0.62434657	0.001	-9.286203273	Down	1.08E-06	1.99E-06	AAEL011921-PA (Fragment) OS=Aedes aegypti GN=AAEL011921 PE=4 SV=1
CCG027833.1	363	18	0	2.13073064	0.001	-11.05713251	Down	4.86E-07	9.07E-07	AAEL017144-PA OS=Aedes aegypti PE=4 SV=1
CCG009213.1	591	18	0	1.30872288	0.001	-10.35394393	Down	4.86E-07	9.07E-07	AAEL009707-PA OS=Aedes aegypti GN=AAEL009707 PE=4 SV=1
CCG012935.1	378	19	0	2.15985439	0.001	-11.07671834	Down	2.18E-07	4.12E-07	AAEL002003-PA OS=Aedes aegypti GN=AAEL002003 PE=4 SV=1
CCG018645.1	552	19	0	1.47903072	0.001	-10.5304363	Down	2.18E-07	4.12E-07	AAEL013917-PA OS=Aedes aegypti GN=AAEL013917 PE=4 SV=1
CCG011150.1	702	19	0	1.16299852	0.001	-10.18363354	Down	2.18E-07	4.12E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015444 PE=4 SV=1

CCG021471.1	588	20	0	1.4615556	0.001	-10.513289	Down	9.77E-08	1.88E-07	AAEL011586-PA OS=Aedes aegypti GN=AAEL011586 PE=4 SV=1
CCG026792.1	978	20	0	0.87872668	0.001	-9.779270686	Down	9.77E-08	1.88E-07	AAEL009463-PA (Fragment) OS=Aedes aegypti GN=AAEL009463 PE=4 SV=1
CCG009628.1	981	20	0	0.87603944	0.001	-9.774852015	Down	9.77E-08	1.88E-07	AAEL004557-PA OS=Aedes aegypti GN=AAEL004557 PE=4 SV=1
CCG023064.1	354	21	0	2.54905205	0.001	-11.31574512	Down	4.38E-08	8.54E-08	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015505 PE=4 SV=1
CCG016431.1	468	21	0	1.92812912	0.001	-10.91298595	Down	4.38E-08	8.54E-08	AAEL002383-PA OS=Aedes aegypti GN=AAEL002383 PE=4 SV=1
CCG020053.1	648	21	0	1.3925377	0.001	-10.44350067	Down	4.38E-08	8.54E-08	-
CCG002196.1	682	21	0	1.323115	0.001	-10.36972274	Down	4.38E-08	8.54E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007192 PE=4 SV=1
CCG014334.1	846	21	0	1.06662462	0.001	-10.05883682	Down	4.38E-08	8.54E-08	Guanine nucleotide exchange factor, putative OS=Pediculus humanus subsp. corporis GN=Phum_PHUM503240 PE=4 SV=1
CCG003729.1	1509	21	0	0.59798835	0.001	-9.223973579	Down	4.38E-08	8.54E-08	AAEL007812-PA OS=Aedes aegypti GN=CYP4H32 PE=3 SV=1
CCG022367.1	273	22	0	3.4627625	0.001	-11.75770772	Down	1.97E-08	3.88E-08	AAEL011229-PA OS=Aedes aegypti GN=AAEL011229 PE=4 SV=1
CCG004628.1	621	22	0	1.52227723	0.001	-10.57201541	Down	1.97E-08	3.88E-08	AAEL008440-PA OS=Aedes aegypti GN=AAEL008440 PE=4 SV=1
CCG020377.1	741	22	0	1.2757546	0.001	-10.31713513	Down	1.97E-08	3.88E-08	AAEL004335-PA OS=Aedes aegypti GN=AAEL004335 PE=4 SV=1
CCG019941.1	783	22	0	1.20732332	0.001	-10.23759637	Down	1.97E-08	3.88E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011970 PE=4 SV=1
CCG011586.1	882	22	0	1.07180744	0.001	-10.06583002	Down	1.97E-08	3.88E-08	-

CCG026493.1	915	22	0	1.03315209	0.001	-10.01283693	Down	1.97E-08	3.88E-08	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG027536.1	618	23	0	1.59919724	0.001	-10.64313217	Down	8.82E-09	1.76E-08	Ecdysis triggering hormone OS=Aedes aegypti PE=2 SV=1
CCG003688.1	654	23	0	1.51116804	0.001	-10.56144838	Down	8.82E-09	1.76E-08	AAEL002275-PA OS=Aedes aegypti GN=AAEL002275 PE=4 SV=1
CCG012487.1	735	23	0	1.34463115	0.001	-10.39299476	Down	8.82E-09	1.76E-08	AGAP013022-PA OS=Anopheles gambiae GN=AgaP_AGAP013022 PE=4 SV=1
CCG003765.1	1407	23	0	0.70241926	0.001	-9.456188589	Down	8.82E-09	1.76E-08	AAEL005004-PA OS=Aedes aegypti GN=AAEL005004 PE=4 SV=1
CCG020393.1	1458	23	0	0.67784904	0.001	-9.404820198	Down	8.82E-09	1.76E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001348 PE=4 SV=1
CCG021507.1	879	24	0	1.17323507	0.001	-10.19627639	Down	3.96E-09	8.00E-09	AAEL010413-PA OS=Aedes aegypti GN=AAEL010413 PE=4 SV=1
CCG018091.1	1215	24	0	0.84878488	0.001	-9.729255149	Down	3.96E-09	8.00E-09	AAEL001053-PA OS=Aedes aegypti GN=AAEL001053 PE=4 SV=1
CCG005599.1	1256	24	0	0.82107773	0.001	-9.681374998	Down	3.96E-09	8.00E-09	AAEL009780-PA OS=Aedes aegypti GN=AAEL009780 PE=3 SV=1
CCG015647.1	1491	24	0	0.69166575	0.001	-9.433931205	Down	3.96E-09	8.00E-09	AAEL009258-PA OS=Aedes aegypti GN=AAEL009258 PE=4 SV=1
CCG009553.1	3315	24	0	0.3110931	0.001	-8.281202592	Down	3.96E-09	8.00E-09	Gamma-aminobutyric acid type B receptor OS=Culex quinquefasciatus GN=CpipJ_CPIJ002466 PE=4 SV=1
CCG017307.1	792	25	0	1.35636789	0.001	-10.40553282	Down	1.78E-09	3.63E-09	AAEL002361-PA OS=Aedes aegypti GN=AAEL002361 PE=4 SV=1
CCG026763.1	489	26	0	2.28468937	0.001	-11.15778231	Down	7.96E-10	1.65E-09	AAEL008473-PA OS=Aedes aegypti GN=AAEL008473 PE=4 SV=1
CCG010494.1	711	26	0	1.57132644	0.001	-10.61776721	Down	7.96E-10	1.65E-09	AAEL009963-PA OS=Aedes aegypti GN=AAEL009963 PE=4 SV=1
CCG010391.1	813	26	0	1.37418585	0.001	-10.42436142	Down	7.96E-10	1.65E-09	-

CCG021870.1	1065	26	0	1.04902638	0.001	-10.03483525	Down	7.96E-10	1.65E-09	AAEL001333-PA OS=Aedes aegypti GN=AAEL001333 PE=4 SV=1
CCG022312.1	1170	26	0	0.95488299	0.001	-9.89918015	Down	7.96E-10	1.65E-09	AAEL001340-PA (Fragment) OS=Aedes aegypti GN=AAEL001340 PE=4 SV=1
CCG004606.1	1302	26	0	0.85807458	0.001	-9.744959231	Down	7.96E-10	1.65E-09	AAEL013008-PA OS=Aedes aegypti GN=AAEL013008 PE=4 SV=1
CCG026537.1	1416	26	0	0.7889923	0.001	-9.623867414	Down	7.96E-10	1.65E-09	AAEL002713-PA (Fragment) OS=Aedes aegypti GN=AAEL002713 PE=4 SV=1
CCG027160.1	2049	26	0	0.54524797	0.001	-9.090768695	Down	7.96E-10	1.65E-09	AGAP002293-PA OS=Anopheles gambiae GN=AGAP002293 PE=4 SV=3
CCG012275.1	468	27	0	2.47902315	0.001	-11.27555603	Down	3.57E-10	7.49E-10	AAEL005222-PA OS=Aedes aegypti GN=AAEL005222 PE=4 SV=1
CCG010435.1	552	27	0	2.1017805	0.001	-11.03739629	Down	3.57E-10	7.50E-10	AAEL012690-PA OS=Aedes aegypti GN=AAEL012690 PE=4 SV=1
CCG020025.1	219	28	0	5.49384735	0.001	-12.42360111	Down	1.60E-10	3.41E-10	Uncharacterized protein OS=Anopheles darlingi GN=AND_08788 PE=4 SV=1
CCG004366.1	285	28	0	4.22158796	0.001	-12.04357006	Down	1.60E-10	3.41E-10	AAEL002878-PA OS=Aedes aegypti GN=AAEL002878 PE=4 SV=1
CCG025446.1	720	28	0	1.67104523	0.001	-10.70653507	Down	1.60E-10	3.41E-10	AAEL003644-PA OS=Aedes aegypti GN=AAEL003644 PE=4 SV=1
CCG005203.1	771	28	0	1.56050917	0.001	-10.60780112	Down	1.60E-10	3.41E-10	AAEL006429-PA OS=Aedes aegypti GN=AAEL006429 PE=3 SV=1
CCG027834.1	363	29	0	3.43284381	0.001	-11.7451885	Down	7.19E-11	1.54E-10	AAEL017144-PA OS=Aedes aegypti PE=4 SV=1
CCG015893.1	831	29	0	1.49954549	0.001	-10.55030957	Down	7.19E-11	1.54E-10	Uncharacterized protein OS=Anopheles darlingi GN=AND_08032 PE=4 SV=1
CCG007862.1	348	30	0	3.70428747	0.001	-11.85498035	Down	3.23E-11	7.00E-11	AAEL000929-PA OS=Aedes aegypti GN=AAEL000929 PE=3 SV=1
CCG027590.1	795	30	0	1.62149942	0.001	-10.66311279	Down	3.23E-11	7.00E-11	-

CCG025964.1	1743	34	0	0.83819333	0.001	-9.711139233	Down	1.31E-12	3.17E-12	AAEL000873-PA (Fragment) OS=Aedes aegypti GN=AAEL000873 PE=3 SV=1
CCG018700.1	897	35	0	1.67663401	0.001	-10.71135209	Down	5.86E-13	1.50E-12	-
CCG008724.1	714	38	0	2.28690464	0.001	-11.1591805	Down	5.29E-14	1.78E-13	AAEL013073-PA OS=Aedes aegypti GN=AAEL013073 PE=3 SV=1
CCG016598.1	294	39	0	5.70006684	0.001	-12.47676312	Down	2.37E-14	8.28E-14	AAEL007735-PA OS=Aedes aegypti GN=AAEL007735 PE=4 SV=1
CCG024954.1	744	39	0	2.25244577	0.001	-11.13727665	Down	2.37E-14	8.27E-14	AAEL013325-PA OS=Aedes aegypti GN=AAEL013325 PE=4 SV=1
CCG015924.1	831	39	0	2.01663014	0.001	-10.9777308	Down	2.37E-14	8.28E-14	Uncharacterized protein OS=Anopheles darlingi GN=AND_08032 PE=4 SV=1
CCG006451.1	561	40	0	3.06379569	0.001	-11.58110438	Down	1.06E-14	3.80E-14	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013022 PE=4 SV=1
CCG011880.1	477	41	0	3.69341534	0.001	-11.85073979	Down	4.78E-15	1.72E-14	-
CCG003041.1	1035	41	0	1.70218272	0.001	-10.7331702	Down	4.78E-15	1.72E-14	AAEL013496-PA OS=Aedes aegypti GN=PPO8 PE=4 SV=1
CCG002045.1	1116	42	0	1.61714055	0.001	-10.65922936	Down	2.14E-15	7.77E-15	AAEL009949-PA OS=Aedes aegypti GN=AAEL009949 PE=3 SV=1
CCG000300.1	1719	42	0	1.04987135	0.001	-10.03599684	Down	2.14E-15	7.76E-15	AAEL009860-PA OS=Aedes aegypti GN=AAEL009860 PE=4 SV=1
CCG010540.1	510	43	0	3.62293841	0.001	-11.82294456	Down	9.61E-16	3.50E-15	AGAP013237-PA OS=Anopheles gambiae GN=AgaP_AGAP013237 PE=4 SV=1
CCG027254.1	1074	43	0	1.72038975	0.001	-10.74851972	Down	9.61E-16	3.50E-15	AAEL009514-PA OS=Aedes aegypti GN=AAEL009514 PE=4 SV=1
CCG011477.1	816	44	0	2.31699549	0.001	-11.17803952	Down	4.31E-16	1.61E-15	AAEL000725-PA OS=Aedes aegypti GN=AAEL000725 PE=4 SV=1
CCG009060.1	969	44	0	1.9511541	0.001	-10.93011201	Down	4.31E-16	1.61E-15	AAEL002689-PA OS=Aedes aegypti GN=AAEL002689 PE=4 SV=1
CCG016082.1	252	45	0	7.67316689	0.001	-12.90560642	Down	1.93E-16	7.26E-16	-
CCG014364.1	1161	45	0	1.66549359	0.001	-10.70173409	Down	1.93E-16	7.26E-16	Lipase OS=Aedes aegypti GN=AAEL004925 PE=3 SV=1

CCG004591.1	2694	45	0	0.71775726	0.001	-9.487352207	Down	1.93E-16	7.26E-16	AAEL005240-PA OS=Aedes aegypti GN=AAEL005240 PE=4 SV=1
CCG026638.1	1104	46	0	1.79040561	0.001	-10.80607075	Down	8.68E-17	3.27E-16	-
CCG007555.1	1098	47	0	1.83932379	0.001	-10.84495976	Down	3.89E-17	1.47E-16	AAEL004705-PA OS=Aedes aegypti GN=AAEL004705 PE=4 SV=1
CCG025751.1	1392	50	0	1.54345311	0.001	-10.59194594	Down	3.51E-18	1.34E-17	Uncharacterized protein OS=Anopheles darlingi GN=AND_13111 PE=3 SV=1
CCG011468.3	2007	50	0	1.07049663	0.001	-10.06406453	Down	3.51E-18	1.34E-17	AAEL010689-PA OS=Aedes aegypti GN=AAEL010689 PE=4 SV=1
CCG011043.1	285	51	0	7.68932093	0.001	-12.90864048	Down	1.58E-18	6.06E-18	AAEL002878-PA OS=Aedes aegypti GN=AAEL002878 PE=4 SV=1
CCG018644.1	708	51	0	3.09527749	0.001	-11.59585304	Down	1.58E-18	6.06E-18	AAEL013918-PA OS=Aedes aegypti GN=AAEL013918 PE=4 SV=1
CCG027147.1	465	52	0	4.80521763	0.001	-12.23038606	Down	7.07E-19	2.73E-18	-
CCG019735.1	1365	52	0	1.63694227	0.001	-10.67678773	Down	7.07E-19	2.73E-18	GD15416 OS=Drosophila simulans GN=GD15416 PE=3 SV=1
CCG017202.1	420	53	0	5.42237127	0.001	-12.40470818	Down	3.17E-19	1.23E-18	AAEL012573-PA OS=Aedes aegypti GN=AAEL012573 PE=4 SV=1
CCG025163.3	363	54	0	6.39219193	0.001	-12.64209501	Down	1.42E-19	5.53E-19	NADH dehydrogenase [ubiquinone] 1 subunit C2 OS=Aedes aegypti GN=AAEL005046 PE=3 SV=1
CCG018289.1	1830	54	0	1.26795938	0.001	-10.30829282	Down	1.42E-19	5.53E-19	Malic enzyme OS=Aedes aegypti GN=AAEL001073 PE=3 SV=1
CCG024212.1	555	55	0	4.25826199	0.001	-12.056049	Down	6.38E-20	2.49E-19	AAEL007754-PA OS=Aedes aegypti GN=AAEL007754 PE=4 SV=1
CCG027815.1	246	56	0	9.7817282	0.001	-13.25587366	Down	2.86E-20	1.12E-19	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011592 PE=4 SV=1
CCG003754.1	540	56	0	4.45612063	0.001	-12.12157257	Down	2.86E-20	1.12E-19	AAEL014879-PA OS=Aedes aegypti GN=AAEL014879 PE=4 SV=1
CCG005716.1	432	57	0	5.66961776	0.001	-12.46903576	Down	1.28E-20	5.04E-20	AAEL004001-PA OS=Aedes aegypti GN=AAEL004001 PE=4 SV=1

CCG012993.1	432	58	0	5.76908474	0.001	-12.49412674	Down	5.76E-21	2.27E-20	Protein disulfide isomerase-like PDI-H OS=Physcomitrella patens PE=2 SV=1
CCG027395.1	327	59	0	7.75294906	0.001	-12.92052947	Down	2.58E-21	1.02E-20	AAEL008308-PA OS=Aedes aegypti GN=AAEL008308 PE=4 SV=1
CCG017067.1	690	61	0	3.79877364	0.001	-11.89131803	Down	5.20E-22	2.06E-21	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015444 PE=4 SV=1
CCG004738.1	1851	62	0	1.43928879	0.001	-10.49114038	Down	2.33E-22	9.29E-22	Hyperpolarization activated cyclic nucleotide-gated potassium channel OS=Culex quinquefasciatus GN=CpipJ_CPIJ001483 PE=4 SV=1
CCG009550.1	843	65	0	3.31320611	0.001	-11.69401224	Down	2.11E-23	8.46E-23	AAEL007706-PA OS=Aedes aegypti GN=AAEL007706 PE=4 SV=1
CCG001524.1	651	66	0	4.35637862	0.001	-12.08891363	Down	9.45E-24	3.80E-23	-
CCG004505.1	1674	66	0	1.69414724	0.001	-10.72634355	Down	9.45E-24	3.80E-23	AAEL007188-PA (Fragment) OS=Aedes aegypti GN=AAEL007188 PE=4 SV=1
CCG000700.1	609	67	0	4.72737639	0.001	-12.20682402	Down	4.24E-24	1.71E-23	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014666 PE=4 SV=1
CCG020247.1	1302	68	0	2.24419505	0.001	-11.13198235	Down	1.90E-24	7.69E-24	Uncharacterized protein OS=Anopheles darlingi GN=AND_15514 PE=4 SV=1
CCG017298.1	471	70	0	6.38616013	0.001	-12.64073301	Down	3.83E-25	1.56E-24	-
CCG000755.1	642	72	0	4.81903566	0.001	-12.23452876	Down	7.70E-26	3.15E-25	AAEL013737-PA OS=Aedes aegypti GN=AAEL013737 PE=4 SV=1
CCG026297.1	1476	72	0	2.09608462	0.001	-11.03348124	Down	7.70E-26	3.15E-25	-
CCG012388.1	1518	73	0	2.06639699	0.001	-11.01290173	Down	3.45E-26	1.42E-25	AAEL001658-PA OS=Aedes aegypti GN=AAEL001658 PE=4 SV=1
CCG019807.1	1611	73	0	1.94710778	0.001	-10.92711703	Down	3.45E-26	1.42E-25	AAEL007121-PA OS=Aedes aegypti GN=AAEL007121 PE=4 SV=1
CCG011522.1	486	74	0	6.54271679	0.001	-12.67567411	Down	1.55E-26	6.38E-26	-

CCG021755.1	1215	74	0	2.61708672	0.001	-11.35374601	Down	1.55E-26	6.38E-26	AAEL002274-PA (Fragment) OS=Aedes aegypti GN=AAEL002274 PE=4 SV=1
CCG010538.1	1290	74	0	2.46493051	0.001	-11.26733126	Down	1.55E-26	6.39E-26	AAEL006399-PA OS=Aedes aegypti GN=AAEL006399 PE=4 SV=1
CCG004014.1	678	75	0	4.75328923	0.001	-12.21471047	Down	6.95E-27	2.87E-26	AAEL002657-PA OS=Aedes aegypti GN=AAEL002657 PE=4 SV=1
CCG024050.1	1725	76	0	1.89315932	0.001	-10.88658011	Down	3.12E-27	1.29E-26	AAEL001658-PA OS=Aedes aegypti GN=AAEL001658 PE=4 SV=1
CCG000598.1	1851	77	0	1.78750382	0.001	-10.80373061	Down	1.40E-27	5.80E-27	Hyperpolarization activated cyclic nucleotide-gated potassium channel OS=Culex quinquefasciatus GN=CpipJ_CPIJ001482 PE=4
CCG020425.1	375	78	0	8.9377048	0.001	-13.12568868	Down	6.28E-28	2.61E-27	-
CCG022406.1	402	79	0	8.44430108	0.001	-13.0437623	Down	2.82E-28	1.17E-27	-
CCG017949.1	897	80	0	3.83230632	0.001	-11.90399717	Down	1.26E-28	5.28E-28	-
CCG023186.1	651	84	0	5.54448189	0.001	-12.43683694	Down	5.11E-30	2.15E-29	-
CCG003636.1	756	85	0	4.83125323	0.001	-12.23818176	Down	2.29E-30	9.69E-30	AAEL009485-PA OS=Aedes aegypti GN=AAEL009485 PE=4 SV=1
CCG017732.1	606	87	0	6.1689223	0.001	-12.59080276	Down	4.62E-31	1.96E-30	AGAP004947-PA OS=Anopheles gambiae GN=AGAP004947 PE=4 SV=1
CCG002324.1	1251	87	0	2.98830289	0.001	-11.54511067	Down	4.62E-31	1.96E-30	AAEL006924-PA OS=Aedes aegypti GN=AAEL006924 PE=4 SV=1
CCG006739.1	561	90	0	6.89354031	0.001	-12.75102938	Down	4.17E-32	1.78E-31	AAEL003217-PA OS=Aedes aegypti GN=AAEL003217 PE=4 SV=1
CCG004986.1	1638	90	0	2.36097443	0.001	-11.2051667	Down	4.17E-32	1.78E-31	AAEL000108-PA OS=Aedes aegypti GN=AAEL000108 PE=4 SV=1
CCG014398.1	1908	98	0	2.20704088	0.001	-11.10789763	Down	6.84E-35	2.96E-34	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018057 PE=4 SV=1
CCG014950.1	1719	99	0	2.47469676	0.001	-11.27303604	Down	3.07E-35	1.33E-34	AAEL009860-PA OS=Aedes aegypti GN=AAEL009860 PE=4 SV=1
CCG006211.1	1806	102	0	2.42686209	0.001	-11.24487641	Down	2.77E-36	1.21E-35	AAEL003976-PA OS=Aedes aegypti GN=AAEL003976 PE=4 SV=1

CCG008090.1	711	103	0	6.22487013	0.001	-12.60382802	Down	1.24E-36	5.44E-36	AAEL009963-PA OS=Aedes aegypti GN=AAEL009963 PE=4 SV=1
CCG002150.1	2247	106	0	2.02705468	0.001	-10.98516929	Down	1.12E-37	4.94E-37	AAEL000382-PA OS=Aedes aegypti GN=AAEL000382 PE=4 SV=1
CCG021138.1	1368	107	0	3.36093684	0.001	-11.71464772	Down	5.03E-38	2.22E-37	GD15416 OS=Drosophila simulans GN=GD15416 PE=3 SV=1
CCG003365.1	681	108	0	6.81458346	0.001	-12.73440976	Down	2.26E-38	9.99E-38	AAEL015481-PA OS=Aedes aegypti GN=AAEL015481 PE=4 SV=1
CCG014194.1	1512	112	0	3.1829433	0.001	-11.63614574	Down	9.14E-40	4.07E-39	AAEL007411-PA OS=Aedes aegypti GN=AAEL007411 PE=4 SV=1
CCG023820.1	627	114	0	7.81267902	0.001	-12.93160163	Down	1.84E-40	8.22E-40	AAEL014067-PA OS=Aedes aegypti GN=AAEL014067 PE=4 SV=1
CCG018151.1	561	116	0	8.88500751	0.001	-13.11715728	Down	3.70E-41	1.66E-40	AAEL003576-PA OS=Aedes aegypti GN=AAEL003576 PE=4 SV=1
CCG006969.1	876	117	0	5.73910839	0.001	-12.48661091	Down	1.66E-41	7.46E-41	AAEL010658-PA OS=Aedes aegypti GN=AAEL010658 PE=4 SV=1
CCG002697.1	1395	121	0	3.72712393	0.001	-11.86384708	Down	6.72E-43	3.03E-42	AAEL005630-PA OS=Aedes aegypti GN=AAEL005630 PE=4 SV=1
CCG002783.1	1245	122	0	4.21068885	0.001	-12.03984056	Down	3.02E-43	1.36E-42	AAEL005714-PA (Fragment) OS=Aedes aegypti GN=AAEL005714 PE=4 SV=1
CCG004579.1	1638	122	0	3.200432	0.001	-11.64405094	Down	3.02E-43	1.36E-42	AAEL000108-PA OS=Aedes aegypti GN=AAEL000108 PE=4 SV=1
CCG001526.1	687	123	0	7.69327126	0.001	-12.90938146	Down	1.35E-43	6.13E-43	-
CCG002603.1	843	126	0	6.42252261	0.001	-12.64892435	Down	1.22E-44	5.56E-44	AAEL007706-PA OS=Aedes aegypti GN=AAEL007706 PE=4 SV=1
CCG002701.1	930	128	0	5.91411401	0.001	-12.52994634	Down	2.46E-45	1.12E-44	AAEL000694-PA OS=Aedes aegypti GN=AAEL000694 PE=4 SV=1
CCG027361.1	489	129	0	11.3355742	0.001	-13.46856985	Down	1.10E-45	5.05E-45	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC005836 PE=4 SV=1
CCG026915.1	1656	129	0	3.34728005	0.001	-11.70877354	Down	1.10E-45	5.05E-45	AAEL002674-PA OS=Aedes aegypti GN=AAEL002674 PE=4 SV=1

CCG023461.1	1629	130	0	3.42913781	0.001	-11.74363017	Down	4.95E-46	2.27E-45	AAEL002674-PA OS=Aedes aegypti GN=AAEL002674 PE=4 SV=1
CCG026162.1	1239	132	0	4.5778894	0.001	-12.16046689	Down	9.95E-47	4.58E-46	Vitelline envelope protein alpha OS=Oncorhynchus mykiss PE=2 SV=1
CCG021524.1	477	133	0	11.981079	0.001	-13.54847023	Down	4.46E-47	2.06E-46	-
CCG001041.1	696	134	0	8.27290868	0.001	-13.01417894	Down	2.00E-47	9.24E-47	-
CCG025671.1	1509	139	0	3.95811339	0.001	-11.95059723	Down	3.64E-49	1.69E-48	AAEL007411-PA OS=Aedes aegypti GN=AAEL007411 PE=4 SV=1
CCG021915.1	1647	139	0	3.62646819	0.001	-11.82434948	Down	3.64E-49	1.69E-48	AAEL002405-PA OS=Aedes aegypti GN=AAEL002405 PE=4 SV=1
CCG008141.1	2343	139	0	2.54920747	0.001	-11.31583308	Down	3.64E-49	1.69E-48	AAEL000382-PA OS=Aedes aegypti GN=AAEL000382 PE=4 SV=1
CCG023518.1	948	141	0	6.39106812	0.001	-12.64184135	Down	7.32E-50	3.41E-49	AAEL005907-PA OS=Aedes aegypti GN=AAEL005907 PE=4 SV=1
CCG023349.1	657	143	0	9.35262108	0.001	-13.19115502	Down	1.47E-50	6.89E-50	AAEL013073-PA OS=Aedes aegypti GN=AAEL013073 PE=3 SV=1
CCG005701.1	642	145	0	9.70500237	0.001	-13.24451285	Down	2.97E-51	1.39E-50	AAEL002052-PA OS=Aedes aegypti GN=AAEL002052 PE=4 SV=1
CCG020854.1	720	148	0	8.83266767	0.001	-13.10863352	Down	2.68E-52	1.26E-51	-
CCG012907.1	1113	148	0	5.7138551	0.001	-12.48024873	Down	2.68E-52	1.26E-51	AAEL002270-PA OS=Aedes aegypti GN=AAEL002270 PE=4 SV=1
CCG001249.1	1248	151	0	5.19906244	0.001	-12.34403577	Down	2.42E-53	1.14E-52	AAEL006924-PA OS=Aedes aegypti GN=AAEL006924 PE=4 SV=1
CCG022067.1	1479	156	0	4.53230466	0.001	-12.14602913	Down	4.39E-55	2.08E-54	AAEL000542-PA OS=Aedes aegypti GN=AAEL000542 PE=4 SV=1
CCG001562.1	588	160	0	11.6924448	0.001	-13.513289	Down	1.78E-56	8.48E-56	-
CCG026163.1	1458	160	0	4.71547156	0.001	-12.20318634	Down	1.78E-56	8.48E-56	Putative uncharacterized protein OS=Harpegnathos saltator GN=EAI_02652 PE=4 SV=1
CCG023341.1	369	167	0	19.4470073	0.001	-14.24726053	Down	6.50E-59	3.12E-58	AAEL003177-PA OS=Aedes aegypti GN=AAEL003177 PE=4 SV=1
CCG006177.1	594	167	0	12.0807166	0.001	-13.56041842	Down	6.50E-59	3.12E-58	-
CCG017154.1	360	170	0	20.2912636	0.001	-14.30857109	Down	5.87E-60	2.83E-59	AAEL012715-PA OS=Aedes aegypti GN=AAEL012715 PE=4 SV=1

CCG020012.1	384	178	0	19.9182624	0.001	-14.28180418	Down	9.62E-63	4.68E-62	-
CCG027782.1	1623	184	0	4.87149179	0.001	-12.25014792	Down	7.84E-65	3.83E-64	Putative protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009617 PE=4 SV=1
CCG013013.1	1173	190	0	6.96014457	0.001	-12.76490156	Down	6.39E-67	3.14E-66	AAEL009564-PA OS=Aedes aegypti GN=AAEL009564 PE=4 SV=1
CCG000021.1	801	191	0	10.2462164	0.001	-13.32280364	Down	2.87E-67	1.41E-66	-
CCG001024.1	699	195	0	11.987265	0.001	-13.54921491	Down	1.16E-68	5.73E-68	AAEL001452-PA OS=Aedes aegypti GN=AAEL001452 PE=4 SV=1
CCG019017.1	513	196	0	16.4172865	0.001	-14.00292807	Down	5.21E-69	2.57E-68	-
CCG006467.1	702	206	0	12.6093523	0.001	-13.62220655	Down	1.72E-72	8.57E-72	AAEL005352-PA OS=Aedes aegypti GN=AAEL005352 PE=4 SV=1
CCG000244.1	1674	211	0	5.4161374	0.001	-12.40304862	Down	3.12E-74	1.57E-73	AAEL007188-PA (Fragment) OS=Aedes aegypti GN=AAEL007188 PE=4 SV=1
CCG017459.1	615	212	0	14.8123313	0.001	-13.8545111	Down	1.40E-74	7.02E-74	AAEL010160-PA OS=Aedes aegypti GN=AAEL010160 PE=4 SV=1
CCG022757.1	1314	215	0	7.03081655	0.001	-12.77947654	Down	1.26E-75	6.36E-75	-
CCG011441.1	1881	219	0	5.00285586	0.001	-12.28853617	Down	5.12E-77	2.58E-76	AAEL015536-PA OS=Aedes aegypti GN=AAEL015536 PE=4 SV=1
CCG023615.1	1503	223	0	6.37541638	0.001	-12.63830385	Down	2.07E-78	1.05E-77	AAEL000542-PA OS=Aedes aegypti GN=AAEL000542 PE=4 SV=1
CCG013078.1	333	227	0	29.291681	0.001	-14.83820337	Down	8.40E-80	4.26E-79	-
CCG014856.1	873	239	0	11.7637647	0.001	-13.52206221	Down	5.58E-84	2.86E-83	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011620 PE=4 SV=1
CCG022597.1	477	240	0	21.6199923	0.001	-14.40007839	Down	2.50E-84	1.28E-83	AAEL001115-PA OS=Aedes aegypti GN=AAEL001115 PE=4 SV=1
CCG021991.1	273	242	0	38.0903875	0.001	-15.21713934	Down	5.04E-85	2.58E-84	-
CCG014875.1	600	243	0	17.4027425	0.001	-14.08702706	Down	2.26E-85	1.16E-84	AAEL009046-PA OS=Aedes aegypti GN=AAEL009046 PE=4 SV=1
CCG014755.1	2616	248	0	4.0735834	0.001	-11.99208273	Down	4.10E-87	2.12E-86	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011070 PE=4 SV=1

CCG005487.1	1434	249	0	7.46127191	0.001	-12.86520587	Down	1.84E-87	9.52E-87	AAEL005004-PA OS=Aedes aegypti GN=AAEL005004 PE=4 SV=1
CCG008102.1	594	250	0	18.0849051	0.001	-14.14249841	Down	8.26E-88	4.27E-87	-
CCG028482.1	1056	252	0	10.2541412	0.001	-13.32391905	Down	1.66E-88	8.61E-88	AAEL010114-PA OS=Aedes aegypti GN=AAEL010114 PE=4 SV=1
CCG023069.1	336	257	0	32.8667315	0.001	-15.00434037	Down	3.02E-90	1.57E-89	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010777 PE=4 SV=1
CCG017948.1	516	259	0	21.568142	0.001	-14.39661428	Down	6.08E-91	3.16E-90	-
CCG017204.1	600	261	0	18.6918346	0.001	-14.19012055	Down	1.22E-91	6.38E-91	AAEL009040-PA OS=Aedes aegypti GN=AAEL009046 PE=4 SV=1
CCG027480.1	696	264	0	16.2988649	0.001	-13.99248387	Down	1.10E-92	5.77E-92	Uncharacterized protein OS=Anopheles darlingi GN=AND_17586 PE=4 SV=1
CCG007804.2	5748	267	0	1.99598454	0.001	-10.96288483	Down	9.97E-94	5.21E-93	AAEL005109-PA OS=Aedes aegypti GN=AAEL005109 PE=4 SV=1
CCG005118.1	945	269	0	12.2315964	0.001	-13.57832509	Down	2.01E-94	1.05E-93	AAEL005027-PA OS=Aedes aegypti GN=AAEL003027 PE=4 SV=1
CCG005751.1	720	279	0	16.6507722	0.001	-14.02330146	Down	6.62E-98	3.49E-97	AAEL009697-PA OS=Aedes aegypti GN=AAEL009697 PE=4 SV=1
CCG016572.1	615	282	0	19.7031954	0.001	-14.266142	Down	5.98E-99	3.16E-98	AAEL010160-PA OS=Aedes aegypti GN=AAEL010160 PE=4 SV=1
CCG000293.1	795	287	0	15.5123444	0.001	-13.92112912	Down	1.09E-100	5.76E-100	-
CCG005290.1	357	288	0	34.6646599	0.001	-15.08117798	Down	4.87E-101	2.58E-100	Cuticle protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003474 PE=4 SV=1
CCG017925.1	1095	290	0	11.3801124	0.001	-13.47422718	Down	9.80E-102	5.20E-101	-
CCG013637.1	1293	304	0	10.1027064	0.001	-13.3024542	Down	1.31E-106	7.00E-106	AAEL004939-PA OS=Aedes aegypti GN=AAEL004939 PE=3 SV=1
CCG018730.1	1572	312	0	8.52834427	0.001	-13.05804996	Down	2.15E-109	1.15E-108	AAEL014288-PA OS=Aedes aegypti GN=AAEL014288 PE=4 SV=1
CCG006934.1	669	318	0	20.4250756	0.001	-14.3180538	Down	1.75E-111	9.43E-111	AAEL011558-PA OS=Aedes aegypti GN=AAEL011558 PE=3 SV=1
CCG016441.1	441	323	0	31.4721639	0.001	-14.94178876	Down	3.18E-113	1.72E-112	-

CCG017614.1	345	329	0	40.9769353	0.001	-15.32252447	Down	2.59E-115	1.40E-114	AAEL013751-PA OS=Aedes aegypti GN=AAEL013751 PE=4 SV=1
CCG016799.1	702	330	0	20.1994479	0.001	-14.30202824	Down	1.16E-115	6.31E-115	AAEL005552-PA OS=Aedes aegypti GN=AAEL005552 PE=4 SV=1
CCG011247.1	819	344	0	18.0483379	0.001	-14.13957836	Down	1.55E-120	8.51E-120	-
CCG021646.1	2442	346	0	6.08825888	0.001	-12.57181399	Down	3.13E-121	1.71E-120	AAEL005595-PA OS=Aedes aegypti GN=AAEL005595 PE=4 SV=1
CCG025202.1	717	347	0	20.7956735	0.001	-14.34399579	Down	1.40E-121	7.69E-121	Uncharacterized protein OS=Anopheles darlingi GN=AND_02268 PE=4 SV=1
CCG019111.1	1839	367	0	8.57525427	0.001	-13.06596373	Down	1.53E-128	8.47E-128	Malic enzyme OS=Aedes aegypti GN=AAEL001073 PE=3 SV=1
CCG026409.1	924	368	0	17.1134874	0.001	-14.06284616	Down	6.86E-129	3.80E-128	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013371 PE=4 SV=1
CCG022783.1	1959	369	0	8.09383975	0.001	-12.98260857	Down	3.08E-129	1.71E-128	AAEL011020-PA OS=Aedes aegypti GN=AAEL011020 PE=4 SV=1
CCG027992.1	855	372	0	18.6956038	0.001	-14.19041145	Down	2.78E-130	1.54E-129	AAEL006418-PA OS=Aedes aegypti GN=AAEL006418 PE=3 SV=1
CCG000615.1	741	385	0	22.3257056	0.001	-14.44641815	Down	8.27E-135	4.62E-134	Serine protease inhibitor dipetalogastin OS=Culex quinquefasciatus GN=CpipJ_CPIJ010521 PE=4 SV=1
CCG026431.1	774	416	0	23.0948444	0.001	-14.49528321	Down	1.33E-145	7.54E-145	Superoxide dismutase OS=Aedes aegypti GN=MNSOD2 PE=3 SV=1
CCG012966.1	1131	420	0	15.9569306	0.001	-13.96189555	Down	5.40E-147	3.06E-146	AAEL009451-PA OS=Aedes aegypti GN=AAEL009451 PE=3 SV=1
CCG011556.1	591	445	0	32.3545379	0.001	-14.98168045	Down	1.07E-155	6.11E-155	AGAP006829-PA OS=Anopheles gambiae GN=CPR59 PE=4 SV=4
CCG007146.1	1488	497	0	14.3521224	0.001	-13.80897648	Down	8.42E-174	4.90E-173	AAEL006017-PA OS=Aedes aegypti GN=AAEL006017 PE=4 SV=1
CCG010349.1	306	509	0	71.4758004	0.001	-16.12516725	Down	5.60E-178	3.27E-177	AAEL003177-PA OS=Aedes aegypti GN=AAEL003177 PE=4 SV=1
CCG019784.1	918	518	0	24.2465387	0.001	-14.56549119	Down	4.12E-181	2.41E-180	-
CCG028082.1	384	590	0	66.0212068	0.001	-16.01064189	Down	3.53E-206	2.12E-205	-

CCG020272.1	357	602	0	72.4587682	0.001	-16.14487266	Down	2.35E-210	1.41E-209	Cuticle protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003474 PE=4 SV=1
CCG017455.1	519	616	0	51.0006869	0.001	-15.63822906	Down	3.14E-215	1.89E-214	AAEL010101-PA OS=Aedes aegypti GN=AAEL010161 PE=4 SV=1
CCG027990.1	759	616	0	34.8739875	0.001	-15.08986371	Down	3.14E-215	1.89E-214	AAEL006403-PA OS=Aedes aegypti GN=AAEL006403 PE=3 SV=1
CCG010213.1	606	631	0	44.7424134	0.001	-15.44935546	Down	1.88E-220	1.14E-219	-
CCG019078.1	1584	634	0	17.1987448	0.001	-14.07001566	Down	1.70E-221	1.03E-220	Glutamate dehydrogenase OS=Aedes aegypti GN=AAEL003757 PE=3 SV=1
CCG008372.1	1002	644	0	27.6172745	0.001	-14.75328333	Down	5.60E-225	3.40E-224	AAEL005010-PA OS=Aedes aegypti GN=AAEL005010 PE=4 SV=1
CCG027515.1	171	677	0	170.119943	0.001	-17.37619276	Down	1.82E-236	1.11E-235	AAEL005219-PA OS=Aedes aegypti GN=AAEL005219 PE=4 SV=1
CCG002914.1	2838	721	0	10.9165534	0.001	-13.41422982	Down	8.74E-252	5.38E-251	Putative uncharacterized protein OS=Candida tropicalis (strain ATCC MYA-3404 / T1) GN=CTRG_04951 PE=4 SV=1
CCG020534.1	489	753	0	66.1681189	0.001	-16.01384865	Down	6.32E-263	3.92E-262	AAEL003027-PA OS=Aedes aegypti GN=AAEL003027 PE=4 SV=1
CCG021899.1	1269	759	0	25.7005741	0.001	-14.64951297	Down	5.15E-265	3.20E-264	AAEL010882-PA OS=Aedes aegypti GN=AAEL010882 PE=4 SV=1
CCG026261.2	1881	802	0	18.3209607	0.001	-14.16120754	Down	5.52E-280	3.44E-279	AAEL015536-PA OS=Aedes aegypti GN=AAEL015536 PE=4 SV=1
CCG019693.1	846	828	0	42.0554849	0.001	-15.36000635	Down	4.90E-289	3.07E-288	AAEL010684-PA OS=Aedes aegypti GN=AAEL010684 PE=4 SV=1
CCG013236.1	741	911	0	52.8278384	0.001	-15.68901076	Down	0	0	Serine protease inhibitor dipetalogastin OS=Culex quinquefasciatus GN=CpipJ_CPIJ010521 PE=4 SV=1
CCG000433.1	822	918	0	47.9880978	0.001	-15.55038901	Down	0	0	AAEL013951-PA OS=Aedes aegypti GN=AAEL013951 PE=3 SV=1
CCG024283.1	402	1061	0	113.41017	0.001	-16.7911905	Down	0	0	AAEL003311-PA OS=Aedes aegypti GN=AAEL003311 PE=4 SV=1

CCG011370.1	234	1203	0	220.908507	0.001	-17.75308945	Down	0	0	AAEL006011-PA OS=Aedes aegypti GN=AAEL006011 PE=4 SV=1
CCG001151.1	510	1296	0	109.193679	0.001	-16.73652981	Down	0	0	-
CCG016951.1	423	1422	0	144.451448	0.001	-17.14022514	Down	0	0	-
CCG002949.1	1251	1624	0	55.7816539	0.001	-15.76750309	Down	0	0	AAEL007281-PA OS=Aedes aegypti GN=AAEL007281 PE=4 SV=1
CCG020274.1	381	1719	0	193.871322	0.001	-17.56473989	Down	0	0	Cuticle protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003476 PE=4 SV=1
CCG014380.1	1623	1833	0	48.5295894	0.001	-15.56657703	Down	0	0	Cytosol aminopeptidase OS=Culex quinquefasciatus GN=CpipJ_CPIJ000990 PE=4 SV=1
CCG017832.1	564	1917	0	146.051385	0.001	-17.15611652	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010091 PE=4 SV=1
CCG008430.1	684	2013	0	126.459175	0.001	-16.94831219	Down	0	0	AAEL008779-PA OS=Aedes aegypti GN=AAEL008779 PE=4 SV=1
CCG016570.1	636	2398	0	162.014817	0.001	-17.30576623	Down	0	0	AAEL010164-PA OS=Aedes aegypti GN=AAEL010164 PE=4 SV=1
CCG014935.1	1245	2947	0	101.712296	0.001	-16.63413456	Down	0	0	AAEL002000-PA OS=Aedes aegypti GN=AAEL002000 PE=3 SV=1
CCG026683.1	1407	3202	0	97.7889767	0.001	-16.57738423	Down	0	0	AAEL003308-PA (Fragment) OS=Aedes aegypti GN=AAEL003308 PE=3 SV=1
CCG026789.1	1485	4650	0	134.551694	0.001	-17.03780103	Down	0	0	-
CCG024282.1	636	5918	0	399.834732	0.001	-18.60904427	Down	0	0	-
CCG025834.1	471	12	1	1.09477031	0.074221394	-3.882649297	Down	0.0004866	0.00076472	AAEL002180-PA OS=Aedes aegypti GN=AAEL002180 PE=4 SV=1
CCG012344.1	177	12	1	2.91320235	0.197504389	-3.882649297	Down	0.0004866	0.00076463	-
CCG022944.1	1077	12	1	0.47877142	0.032458939	-3.882649297	Down	0.0004866	0.00076468	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016112 PE=4 SV=1
CCG021308.1	558	13	1	1.00108701	0.062649242	-3.998126514	Down	0.000233	0.00037322	AAEL001290-PA OS=Aedes aegypti GN=AAEL001290 PE=4 SV=1

CCG022613.1	384	13	1	1.45470456	0.091037179	-3.998126514	Down	0.000233	0.00037324	AAEL002022-PA OS=Aedes aegypti GN=AAEL002022 PE=4 SV=1
CCG014571.1	1056	14	1	0.56967451	0.033104429	-4.105041718	Down	0.0001111	0.00018099	AAEL005490-PA OS=Aedes aegypti GN=AAEL005490 PE=4 SV=1
CCG017208.1	1017	14	1	0.59152044	0.03437392	-4.105041718	Down	0.0001111	0.000181	AAEL001491-PA OS=Aedes aegypti GN=AAEL001491 PE=4 SV=1
CCG021678.1	531	15	1	1.21383431	0.065834796	-4.204577392	Down	5.28E-05	8.83E-05	AAEL003069-PA OS=Aedes aegypti GN=AAEL003069 PE=4 SV=1
CCG005618.1	522	15	1	1.23476249	0.066969879	-4.204577392	Down	5.28E-05	8.82E-05	AAEL010040-PA OS=Aedes aegypti GN=AAEL010040 PE=4 SV=1
CCG016938.1	372	15	1	1.73265059	0.093973862	-4.204577392	Down	5.28E-05	8.83E-05	AAEL002191-PA OS=Aedes aegypti GN=AAEL002191 PE=4 SV=1
CCG017549.1	3555	15	1	0.1813069	0.009833552	-4.204577392	Down	5.28E-05	8.83E-05	Carboxypeptidase AT OS=Culex quinquefasciatus GN=CpipJ_CPIJ007169 PE=3 SV=1
CCG028567.1	300	15	1	2.14848673	0.116527589	-4.204577392	Down	5.28E-05	8.83E-05	AAEL005157-PA (Fragment) OS=Aedes aegypti GN=AAEL005157 PE=4 SV=1
CCG017908.1	321	16	1	2.14179363	0.108904289	-4.297686796	Down	2.50E-05	4.29E-05	-
CCG026455.1	1119	16	1	0.61440192	0.031240641	-4.297686796	Down	2.50E-05	4.29E-05	AAEL006817-PA OS=Aedes aegypti GN=AAEL006817 PE=4 SV=1
CCG028306.1	213	17	1	3.42950934	0.164123365	-4.385149637	Down	1.18E-05	2.06E-05	AAEL003475-PA OS=Aedes aegypti GN=AAEL003475 PE=3 SV=1
CCG001156.1	945	17	1	0.77300052	0.036992885	-4.385149637	Down	1.18E-05	2.06E-05	AAEL008205-PA OS=Aedes aegypti GN=AAEL008205 PE=4 SV=1
CCG007564.1	399	17	1	1.8307907	0.087614729	-4.385149637	Down	1.18E-05	2.06E-05	-
CCG015404.1	579	18	1	1.33584667	0.060376989	-4.467611798	Down	5.57E-06	9.89E-06	AAEL004293-PA OS=Aedes aegypti GN=AAEL004293 PE=4 SV=1
CCG014958.1	510	18	1	1.51657887	0.068545641	-4.467611798	Down	5.57E-06	9.89E-06	-
CCG001214.1	753	22	1	1.25542385	0.046425334	-4.757118415	Down	2.69E-07	5.06E-07	AAEL004996-PA OS=Aedes aegypti GN=AAEL004996 PE=3 SV=1
CCG013340.1	1509	22	1	0.62646399	0.023166519	-4.757118415	Down	2.69E-07	5.06E-07	AAEL003775-PA OS=Aedes aegypti GN=CYP325R1 PE=3 SV=1

CCG011467.1	528	26	1	2.1159339	0.066208858	-4.998126514	Down	1.27E-08	2.52E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013074 PE=4 SV=1
CCG014451.1	1728	26	1	0.64653536	0.020230484	-4.998126514	Down	1.27E-08	2.52E-08	AAEL009193-PA OS=Aedes aegypti GN=AAEL009193 PE=3 SV=1
CCG020953.1	1376	26	1	0.81192812	0.025405724	-4.998126514	Down	1.27E-08	2.52E-08	Putative uncharacterized protein GLEAN_08496 OS=Tribolium castaneum GN=GLEAN_08496 PE=4 SV=1
CCG020974.1	750	26	1	1.48961747	0.046611036	-4.998126514	Down	1.27E-08	2.52E-08	AAEL002919-PA OS=Aedes aegypti GN=AAEL002919 PE=4 SV=1
CCG006053.1	513	27	1	2.26156498	0.068144789	-5.052574298	Down	5.87E-09	1.18E-08	AAEL014135-PA OS=Aedes aegypti GN=AAEL014135 PE=4 SV=1
CCG020883.1	1857	27	1	0.62476189	0.018825136	-5.052574298	Down	5.87E-09	1.18E-08	AGAP013374-PA OS=Anopheles gambiae GN=AgaP_AGAP013374 PE=4 SV=1
CCG000149.1	690	27	1	1.6814244	0.050664169	-5.052574298	Down	5.87E-09	1.18E-08	AAEL014993-PA OS=Aedes aegypti GN=AAEL014993 PE=4 SV=1
CCG028271.1	1088	29	1	1.145333	0.032130769	-5.155667791	Down	1.26E-09	2.60E-09	-
CCG016802.2	213	29	1	5.85033945	0.164123365	-5.155667791	Down	1.26E-09	2.60E-09	-
CCG023995.1	207	31	1	6.43508103	0.168880564	-5.251883107	Down	2.70E-10	5.68E-10	AAEL005316-PA OS=Aedes aegypti GN=AAEL005316 PE=4 SV=1
CCG012266.1	328	31	1	4.06116394	0.106580112	-5.251883107	Down	2.70E-10	5.68E-10	-
CCG019299.1	351	32	1	3.91746868	0.09959623	-5.297686796	Down	1.25E-10	2.66E-10	AAEL001801-PA OS=Aedes aegypti GN=AAEL001801 PE=4 SV=1
CCG004896.1	447	33	1	3.17226229	0.078206436	-5.342080916	Down	5.75E-11	1.24E-10	AAEL011685-PA OS=Aedes aegypti GN=AAEL011685 PE=4 SV=1
CCG005780.1	309	33	1	4.58900078	0.113133582	-5.342080916	Down	5.75E-11	1.24E-10	AAEL013161-PA OS=Aedes aegypti GN=AAEL013161 PE=3 SV=1
CCG024787.1	366	34	1	3.99172398	0.095514417	-5.385149637	Down	2.65E-11	5.78E-11	AAEL011906-PA OS=Aedes aegypti GN=AAEL011906 PE=3 SV=1
CCG013146.1	854	35	1	1.7610547	0.04093475	-5.426969813	Down	1.22E-11	2.71E-11	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG021512.1	1068	37	1	1.48865185	0.032732469	-5.507140162	Down	2.59E-12	6.09E-12	AAEL001333-PA OS=Aedes aegypti GN=AAEL001333 PE=4 SV=1

CCG013025.1	319	40	1	5.3880545	0.109587075	-5.619614891	Down	2.51E-13	7.06E-13	-
CCG017840.1	357	45	1	5.4163531	0.097922344	-5.789539893	Down	5.10E-15	1.84E-14	AAEL002878-PA OS=Aedes aegypti GN=AAEL002878 PE=4 SV=1
CCG003575.1	339	45	1	5.70394707	0.10312176	-5.789539893	Down	5.10E-15	1.84E-14	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012243 PE=4 SV=1
CCG021556.1	1170	61	1	2.2403024	0.029878869	-6.228424134	Down	1.83E-20	7.17E-20	AAEL008325-PA OS=Aedes aegypti GN=AAEL008325 PE=4 SV=1
CCG004992.1	561	90	1	6.89354031	0.062314219	-6.789539893	Down	2.13E-30	9.01E-30	AAEL006357-PA OS=Aedes aegypti GN=AAEL006357 PE=4 SV=1
CCG011671.1	441	105	1	10.2308892	0.079270469	-7.011932314	Down	1.49E-35	6.46E-35	-
CCG010282.1	921	106	1	4.9454852	0.037956869	-7.025607251	Down	6.73E-36	2.93E-35	Ams protein transcript variant 1 (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG017393.1	792	123	1	6.67333	0.044139238	-7.240201302	Down	9.38E-42	4.22E-41	AAEL006758-PA OS=Aedes aegypti GN=AAEL006758 PE=4 SV=1
CCG002913.1	1626	135	1	3.56759789	0.021499555	-7.374502393	Down	6.83E-46	3.13E-45	-
CCG021030.1	1260	135	1	4.60390014	0.027744664	-7.374502393	Down	6.83E-46	3.13E-45	Lipase OS=Aedes aegypti GN=AAEL012342 PE=3 SV=1
CCG023716.1	531	139	1	11.2481979	0.065834796	-7.416627869	Down	2.84E-47	1.31E-46	-
CCG009182.1	381	150	1	16.9172184	0.091754007	-7.526505487	Down	4.54E-51	2.12E-50	AAEL000680-PA OS=Aedes aegypti GN=AAEL000680 PE=4 SV=1
CCG023166.1	522	163	1	13.4177524	0.066969879	-7.64641495	Down	1.47E-55	6.98E-55	-
CCG027991.1	756	167	1	9.49199164	0.046241107	-7.681391089	Down	6.08E-57	2.91E-56	AAEL006414-PA OS=Aedes aegypti GN=AAEL006414 PE=3 SV=1
CCG002290.1	486	171	1	15.1189807	0.071930611	-7.715539311	Down	2.52E-58	1.21E-57	-
CCG028512.1	816	178	1	9.37329995	0.042841025	-7.773420227	Down	9.59E-61	4.64E-60	AAEL000725-PA OS=Aedes aegypti GN=AAEL000725 PE=4 SV=1
CCG014453.1	1890	203	1	4.61526779	0.018496443	-7.963022713	Down	2.16E-69	1.07E-68	AAEL009204-PA OS=Aedes aegypti GN=AAEL009204 PE=3 SV=1
CCG010212.1	396	207	1	22.4614522	0.088278477	-7.991173754	Down	8.92E-71	4.43E-70	-
CCG005574.1	249	210	1	36.2395352	0.140394686	-8.011932314	Down	8.17E-72	4.07E-71	-
CCG019431.1	999	211	1	9.07568969	0.03499327	-8.018785985	Down	3.68E-72	1.83E-71	AAEL005714-PA (Fragment) OS=Aedes aegypti GN=AAEL005714 PE=4 SV=1

CCG003506.1	228	220	1	41.4620246	0.153325775	-8.07904651	Down	2.82E-75	1.42E-74	-
CCG026509.1	1008	259	1	11.0408346	0.03468083	-8.314495084	Down	8.77E-89	4.55E-88	AAEL010886-PA OS=Aedes aegypti GN=AAEL010886 PE=4 SV=1
CCG012757.1	573	308	1	23.0971697	0.061009209	-8.564473337	Down	9.09E-106	4.85E-105	AAEL008754-PA OS=Aedes aegypti GN=AAEL008754 PE=4 SV=1
CCG026055.1	360	321	1	38.31468	0.097106324	-8.624116283	Down	2.82E-110	1.52E-109	AAEL012715-PA OS=Aedes aegypti GN=AAEL012715 PE=4 SV=1
CCG013939.1	1143	341	1	12.8194921	0.030584669	-8.711314725	Down	3.26E-117	1.78E-116	AAEL014193-PA OS=Aedes aegypti GN=AAEL014193 PE=4 SV=1
CCG001525.1	666	355	1	22.904288	0.052489905	-8.769362011	Down	4.54E-122	2.49E-121	-
CCG004697.1	780	358	1	19.7220064	0.044818304	-8.781502573	Down	4.13E-123	2.27E-122	-
CCG005142.1	1413	448	1	13.6238083	0.024740465	-9.105041718	Down	2.40E-154	1.37E-153	AAEL010365-PA OS=Aedes aegypti GN=AAEL010365 PE=4 SV=1
CCG016891.1	204	455	1	95.8393591	0.171364102	-9.127409531	Down	8.90E-157	5.10E-156	AGAP003937-PC OS=Anopheles gambiae GN=AgaP_AGAP003937 PE=4 SV=1
CCG016995.1	1008	503	1	21.4422386	0.03468083	-9.272101386	Down	1.91E-173	1.11E-172	AAEL010886-PA OS=Aedes aegypti GN=AAEL010886 PE=4 SV=1
CCG010562.1	450	534	1	50.9907517	0.077685059	-9.358382728	Down	3.28E-184	1.93E-183	AAEL005891-PA OS=Aedes aegypti GN=AAEL005891 PE=4 SV=1
CCG019955.1	420	813	1	83.1771291	0.083233992	-9.964798338	Down	3.67E-281	2.29E-280	AAEL012573-PA OS=Aedes aegypti GN=AAEL012573 PE=4 SV=1
CCG026112.1	1161	1123	1	41.5633178	0.030110488	-10.43082901	Down	0	0	AAEL009025-PA OS=Aedes aegypti GN=AAEL009025 PE=4 SV=1
CCG006909.1	1056	1492	1	60.7110266	0.033104429	-10.84071862	Down	0	0	AAEL010114-PA OS=Aedes aegypti GN=AAEL010114 PE=4 SV=1
CCG002228.1	1188	1601	1	57.9078663	0.029426159	-10.94244439	Down	0	0	AAEL002715-PA OS=Aedes aegypti GN=SRPN22 PE=3 SV=1
CCG023196.1	1245	1699	1	58.6390194	0.028078937	-11.02815693	Down	0	0	AAEL002000-PA OS=Aedes aegypti GN=AAEL002000 PE=3 SV=1
CCG005291.1	549	1722	1	134.779386	0.063676278	-11.04755622	Down	0	0	Cuicle protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003477 PE=4 SV=1
CCG017756.1	1269	1748	1	59.189201	0.027547893	-11.06917627	Down	0	0	AAEL010882-PA OS=Aedes aegypti GN=AAEL010882 PE=4 SV=1

CCG006118.1	1608	7043	1	188.206369	0.021740222	-13.07966116	Down	0	0	AAEL006975-PA (Fragment) OS=Aedes aegypti GN=AAEL006975 PE=4 SV=1
CCG008175.1	239	14	2	2.51705558	0.292537881	-3.105041718	Down	0.0005487	0.00085961	-
CCG028426.1	1281	14	2	0.46961459	0.054579667	-3.105041718	Down	0.0005487	0.00085966	Reverse transcriptase OS=Aedes aegypti GN=pol PE=4 SV=1
CCG004223.1	777	14	2	0.77422945	0.089982694	-3.105041718	Down	0.0005487	0.00085982	AGAP003797-PA OS=Anopheles gambiae GN=AGAP003797 PE=4 SV=1
CCG012202.1	1365	14	2	0.44071523	0.051220918	-3.105041718	Down	0.0005487	0.00085972	AAEL014617-PA OS=Aedes aegypti GN=CYP9J28 PE=3 SV=1
CCG016812.1	720	14	2	0.83552262	0.097106324	-3.105041718	Down	0.0005487	0.00085977	AAEL008660-PA (Fragment) OS=Aedes aegypti GN=AAEL008660 PE=4 SV=1
CCG009296.1	2523	15	2	0.2554681	0.027711674	-3.204577392	Down	0.0002753	0.00043902	AGAP002803-PA OS=Anopheles gambiae GN=AgaP_AGAP002803 PE=4 SV=2
CCG005048.1	1058	15	2	0.60921174	0.066083699	-3.204577392	Down	0.0002753	0.00043899	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018804 PE=4 SV=1
CCG023862.1	465	16	2	1.4785285	0.15035818	-3.297686796	Down	0.0001373	0.0002225	-
CCG007278.1	433	16	2	1.5877962	0.161470101	-3.297686796	Down	0.0001373	0.00022249	Uncharacterized protein OS=Anopheles darlingi GN=AND_04255 PE=4 SV=1
CCG014482.1	1130	17	2	0.64644733	0.061873056	-3.385149637	Down	6.81E-05	0.00011305	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG001130.1	513	17	2	1.42394832	0.136289578	-3.385149637	Down	6.81E-05	0.00011305	Putative 67B2 family heat shock protein OS=Aedes albopictus PE=2 SV=1
CCG020294.1	363	17	2	2.01235672	0.192607586	-3.385149637	Down	6.81E-05	0.00011304	AAEL017144-PA OS=Aedes aegypti PE=4 SV=1
CCG007560.1	409	18	2	1.89108856	0.170945119	-3.467611798	Down	3.36E-05	5.69E-05	-
CCG017726.1	690	18	2	1.1209496	0.101328338	-3.467611798	Down	3.36E-05	5.69E-05	AAEL006280-PA OS=Aedes aegypti GN=AAEL006280 PE=2 SV=1

CCG002655.1	405	18	2	1.90976598	0.172633466	-3.467611798	Down	3.36E-05	5.69E-05	AAEL009489-PA OS=Aedes aegypti GN=AAEL009489 PE=4 SV=1
CCG010515.1	906	19	2	0.9011313	0.077170589	-3.54561431	Down	1.65E-05	2.85E-05	AAEL009792-PA OS=Aedes aegypti GN=AAEL009792 PE=4 SV=1
CCG018128.1	468	19	2	1.74449777	0.149394345	-3.54561431	Down	1.65E-05	2.85E-05	AAEL002383-PA OS=Aedes aegypti GN=AAEL002383 PE=4 SV=1
CCG015679.1	423	19	2	1.93008264	0.165287361	-3.54561431	Down	1.65E-05	2.85E-05	AAEL008483-PA OS=Aedes aegypti GN=AAEL008483 PE=4 SV=1
CCG005128.1	456	19	2	1.79040561	0.153325775	-3.54561431	Down	1.65E-05	2.85E-05	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG012917.1	2907	20	2	0.29562941	0.024051102	-3.619614891	Down	8.09E-06	1.43E-05	Putative uncharacterized protein OS=Drosophila melanogaster GN=ORF1 PE=4 SV=1
CCG007260.1	1479	20	2	0.5810647	0.047272856	-3.619614891	Down	8.09E-06	1.43E-05	AAEL011770-PA OS=Aedes aegypti GN=CYP325L1 PE=3 SV=1
CCG019781.1	1077	20	2	0.79795236	0.064917877	-3.619614891	Down	8.09E-06	1.43E-05	Hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG027804.1	1122	20	2	0.76594892	0.062314219	-3.619614891	Down	8.09E-06	1.43E-05	Leucine rich protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000810 PE=4 SV=1
CCG024431.1	2802	21	2	0.32204298	0.024952375	-3.690004219	Down	3.95E-06	7.07E-06	AAEL013366-PA (Fragment) OS=Aedes aegypti GN=AAEL013366 PE=4 SV=1
CCG022928.1	1017	21	2	0.88728066	0.06874784	-3.690004219	Down	3.95E-06	7.07E-06	AAEL001491-PA OS=Aedes aegypti GN=AAEL001491 PE=4 SV=1
CCG016035.1	438	22	2	2.15829717	0.159626835	-3.757118415	Down	1.92E-06	3.49E-06	Uncharacterized protein OS=Anopheles darlingi GN=AND_09803 PE=4 SV=1
CCG021766.1	426	22	2	2.21909428	0.164123365	-3.757118415	Down	1.92E-06	3.49E-06	-
CCG003508.1	543	23	2	1.82008084	0.128759767	-3.821248752	Down	9.30E-07	1.72E-06	AAEL001552-PA OS=Aedes aegypti GN=AAEL001552 PE=4 SV=1
CCG004273.1	894	23	2	1.10548534	0.078206436	-3.821248752	Down	9.30E-07	1.72E-06	AAEL011014-PA OS=Aedes aegypti GN=AAEL011014 PE=4 SV=1
CCG020914.1	1047	23	2	0.94393877	0.066777988	-3.821248752	Down	9.30E-07	1.72E-06	AAEL004031-PA OS=Aedes aegypti GN=AAEL004031 PE=4 SV=1

CCG013052.1	594	24	2	1.73615089	0.117704636	-3.882649297	Down	4.50E-07	8.40E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002324 PE=4 SV=1
CCG023057.1	468	25	2	2.29539181	0.149394345	-3.941542986	Down	2.17E-07	4.10E-07	AAEL005222-PA OS=Aedes aegypti GN=AAEL005222 PE=4 SV=1
CCG015281.1	423	25	2	2.53958242	0.165287361	-3.941542986	Down	2.17E-07	4.10E-07	-
CCG007790.1	309	25	2	3.47651575	0.226267164	-3.941542986	Down	2.17E-07	4.10E-07	-
CCG002956.1	516	26	2	2.16514167	0.135497197	-3.998126514	Down	1.04E-07	2.00E-07	AAEL003442-PA OS=Aedes aegypti GN=AAEL003442 PE=4 SV=1
CCG003292.1	537	26	2	2.08047132	0.130198424	-3.998126514	Down	1.04E-07	2.00E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003408 PE=4 SV=1
CCG023489.1	1569	28	2	0.76682764	0.04456122	-4.105041718	Down	2.39E-08	4.71E-08	AAEL000512-PA OS=Aedes aegypti GN=AAEL000512 PE=4 SV=1
CCG003564.1	942	29	2	1.32284746	0.074221394	-4.155667791	Down	1.14E-08	2.28E-08	AAEL001318-PA OS=Aedes aegypti GN=AAEL001318 PE=4 SV=1
CCG028269.1	993	29	2	1.25490665	0.070409419	-4.155667791	Down	1.14E-08	2.28E-08	AAEL004705-PA OS=Aedes aegypti GN=AAEL004705 PE=4 SV=1
CCG003832.1	600	31	2	2.22010295	0.116527589	-4.251883107	Down	2.59E-09	5.29E-09	AAEL009881-PA OS=Aedes aegypti GN=AAEL009881 PE=4 SV=1
CCG009217.1	510	31	2	2.61188583	0.137091281	-4.251883107	Down	2.59E-09	5.29E-09	-
CCG017743.1	822	32	2	1.67278772	0.085056634	-4.297686796	Down	1.23E-09	2.54E-09	NADH dehydrogenase iron-sulfur protein 3, mitochondrial OS=Culex quinquefasciatus GN=CpipJ_CPIJ013503 PE=3 SV=1
CCG018967.1	324	32	2	4.24392441	0.215791832	-4.297686796	Down	1.23E-09	2.54E-09	-
CCG025476.1	1032	33	2	1.37403221	0.067748598	-4.342080916	Down	5.84E-10	1.22E-09	AAEL002915-PA OS=Aedes aegypti GN=AAEL002915 PE=4 SV=1
CCG010945.1	849	34	2	1.72081387	0.082351653	-4.385149637	Down	2.77E-10	5.82E-10	Mariner Mos1 transposase OS=Acromyrmex echinatio GN=G5I_10332 PE=4 SV=1
CCG001689.1	390	34	2	3.74607943	0.179273214	-4.385149637	Down	2.77E-10	5.82E-10	AAEL014975-PA OS=Aedes aegypti GN=AAEL014975 PE=4 SV=1
CCG008367.1	879	34	2	1.66208302	0.079541017	-4.385149637	Down	2.77E-10	5.82E-10	AAEL010413-PA OS=Aedes aegypti GN=AAEL010413 PE=4 SV=1

CCG027832.1	363	35	2	4.14308736	0.192607586	-4.426969813	Down	1.31E-10	2.79E-10	AAEL017144-PA OS=Aedes aegypti PE=4 SV=1
CCG022401.1	885	36	2	1.74792141	0.079001755	-4.467611798	Down	6.18E-11	1.33E-10	AAEL001043-PA OS=Aedes aegypti GN=AAEL001045 PE=4 SV=1
CCG026197.1	561	37	2	2.83401102	0.124628438	-4.507140162	Down	2.92E-11	6.34E-11	AAEL014103-PA OS=Aedes aegypti GN=AAEL014103 PE=4 SV=1
CCG020912.1	1863	40	2	0.92259226	0.037529014	-4.619614891	Down	3.04E-12	7.12E-12	AAEL004003-PA OS=Aedes aegypti GN=AAEL004003 PE=3 SV=1
CCG003635.1	810	42	2	2.22806031	0.086316733	-4.690004219	Down	6.69E-13	1.69E-12	AAEL009485-PA OS=Aedes aegypti GN=AAEL009485 PE=4 SV=1
CCG027810.1	2544	42	2	0.709406	0.027482922	-4.690004219	Down	6.69E-13	1.69E-12	AAEL010958-PA OS=Aedes aegypti GN=AAEL010958 PE=4 SV=1
CCG020180.1	858	43	2	2.15349486	0.081487825	-4.723951551	Down	3.14E-13	8.62E-13	AAEL014579-PA OS=Aedes aegypti GN=AAEL014579 PE=3 SV=1
CCG011063.1	1863	44	2	1.01485149	0.037529014	-4.757118415	Down	1.47E-13	4.38E-13	AAEL004003-PA OS=Aedes aegypti GN=AAEL004003 PE=3 SV=1
CCG009159.1	1215	51	2	1.80366787	0.057544489	-4.970112138	Down	7.07E-16	2.58E-15	Lipase OS=Aedes aegypti GN=AAEL012349 PE=3 SV=1
CCG024301.1	921	53	2	2.4727426	0.075913739	-5.025607251	Down	1.53E-16	5.75E-16	AAEL006841-PA OS=Aedes aegypti GN=AAEL006841 PE=4 SV=1
CCG005043.1	294	56	2	8.18471135	0.237811407	-5.105041718	Down	1.53E-17	5.83E-17	AAEL007735-PA OS=Aedes aegypti GN=AAEL007735 PE=4 SV=1
CCG023884.1	351	58	2	7.10041199	0.19919246	-5.155667791	Down	3.29E-18	1.26E-17	AAEL001801-PA OS=Aedes aegypti GN=AAEL001801 PE=4 SV=1
CCG008963.1	1212	59	2	2.09176101	0.057686925	-5.180329846	Down	1.53E-18	5.87E-18	AAEL009325-PA OS=Aedes aegypti GN=AAEL009325 PE=4 SV=1
CCG024678.1	1611	64	2	1.70705339	0.043399475	-5.297686796	Down	3.24E-20	1.27E-19	AAEL013415-PA OS=Aedes aegypti GN=AAEL013415 PE=4 SV=1
CCG020434.1	1566	67	2	1.83842415	0.044646586	-5.363775987	Down	3.19E-21	1.26E-20	AAEL003576-PA OS=Aedes aegypti GN=AAEL003576 PE=4 SV=1
CCG000302.1	630	71	2	4.84262088	0.110978656	-5.447433916	Down	1.44E-22	5.74E-22	AAEL000359-PA OS=Aedes aegypti GN=AAEL000359 PE=4 SV=1
CCG006174.1	1104	75	2	2.91913958	0.063330212	-5.526505487	Down	6.48E-24	2.61E-23	AAEL011375-PA OS=Aedes aegypti GN=AAEL011375 PE=3 SV=1

CCG027801.1	1902	78	2	1.76216577	0.036759492	-5.583089015	Down	6.31E-25	2.56E-24	AAEL011020-PA OS=Anopheles aegypti GN=AAEL011020 PE=4 SV=1
CCG027982.1	468	87	2	7.98796349	0.149394345	-5.740630292	Down	5.73E-28	2.38E-27	-
CCG002312.1	855	89	2	4.47287296	0.081773747	-5.773420227	Down	1.20E-28	5.04E-28	-
CCG008063.1	3114	90	2	1.24189984	0.022452329	-5.789539893	Down	5.52E-29	2.32E-28	Uncharacterized protein OS=Anopheles darlingi GN=AND_00371 PE=4 SV=1
CCG009968.1	438	90	2	8.82939752	0.159626835	-5.789539893	Down	5.52E-29	2.31E-28	AAEL010134-PA OS=Anopheles aegypti GN=AAEL010134 PE=4 SV=1
CCG023488.2	2808	91	2	1.3925377	0.024899058	-5.805481436	Down	2.53E-29	1.06E-28	AAEL000512-PA OS=Anopheles aegypti GN=AAEL000512 PE=4 SV=1
CCG025295.1	915	98	2	4.60222294	0.076411534	-5.91239664	Down	1.07E-31	4.54E-31	-
CCG006529.1	771	107	2	5.96337432	0.090682949	-6.039153783	Down	9.31E-35	4.03E-34	AAEL003030-PA OS=Anopheles aegypti GN=AAEL003030 PE=4 SV=1
CCG026290.1	903	108	2	5.13923736	0.07742697	-6.052574298	Down	4.25E-35	1.84E-34	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003706 PE=4 SV=1
CCG006435.1	711	110	2	6.64791956	0.098335518	-6.07904651	Down	8.86E-36	3.86E-35	-
CCG005630.1	654	115	2	7.55584018	0.106906045	-6.143176847	Down	1.76E-37	7.73E-37	AAEL002275-PA OS=Anopheles aegypti GN=AAEL002275 PE=4 SV=1
CCG000255.1	1397	122	2	3.75254662	0.05004764	-6.228424134	Down	7.20E-40	3.21E-39	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004282 PE=4 SV=1
CCG007880.1	516	137	2	11.4086311	0.135497197	-6.395718879	Down	5.41E-45	2.47E-44	AGAP006830-PA OS=Anopheles gambiae GN=CPR58 PE=4 SV=3
CCG010203.1	906	145	2	6.87705466	0.077170589	-6.477595886	Down	9.92E-48	4.58E-47	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003706 PE=4 SV=1
CCG005036.1	1869	148	2	3.40263281	0.037408536	-6.507140162	Down	9.32E-49	4.32E-48	AAEL000919-PA OS=Anopheles aegypti GN=AAEL000919 PE=4 SV=1
CCG027811.1	168	153	2	39.1331512	0.416169962	-6.555074639	Down	1.81E-50	8.44E-50	Uncharacterized protein OS=Anopheles darlingi GN=AND_24739 PE=4 SV=1

CCG007707.1	1875	182	2	4.17092891	0.037288829	-6.805481436	Down	2.03E-60	9.83E-60	Heat shock protein 70 B2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014564 PE=3 SV=1
CCG005673.1	1389	236	2	7.30083324	0.050335892	-7.180329846	Down	5.38E-79	2.73E-78	AAEL005630-PA OS=Aedes aegypti GN=AAEL005630 PE=4 SV=1
CCG014122.1	1872	241	2	5.53189425	0.037348586	-7.210576132	Down	1.02E-80	5.18E-80	AAEL000919-PA OS=Aedes aegypti GN=AAEL000919 PE=4 SV=1
CCG018732.1	1416	257	2	7.79888545	0.049376097	-7.303311345	Down	3.11E-86	1.60E-85	Uncharacterized protein OS=Anopheles darlingi GN=AND_15514 PE=4 SV=1
CCG014435.1	606	272	2	19.2867456	0.115373851	-7.385149637	Down	2.09E-91	1.09E-90	-
CCG007958.1	1248	279	2	9.60621471	0.056022879	-7.421808108	Down	8.02E-94	4.20E-93	AAEL001713-PA OS=Aedes aegypti GN=AAEL001713 PE=4 SV=1
CCG021173.1	2949	280	2	4.07986629	0.023708563	-7.426969813	Down	3.62E-94	1.90E-93	AAEL001340-PA (Fragment) OS=Aedes aegypti GN=AAEL001340 PE=4 SV=1
CCG018620.1	1314	293	2	9.58153139	0.053208945	-7.492443651	Down	1.18E-98	6.24E-98	-
CCG009447.1	876	321	2	15.7457589	0.079813417	-7.624116283	Down	2.53E-108	1.36E-107	AAEL010658-PA OS=Aedes aegypti GN=AAEL010658 PE=4 SV=1
CCG021523.1	426	322	2	32.4794708	0.164123365	-7.628603674	Down	1.14E-108	6.12E-108	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010703 PE=4 SV=1
CCG004722.1	1488	333	2	9.61621077	0.046986931	-7.677065163	Down	1.80E-112	9.74E-112	AAEL006017-PA OS=Aedes aegypti GN=AAEL006017 PE=4 SV=1
CCG026991.1	1923	372	2	8.31239796	0.036358062	-7.836845607	Down	5.95E-126	3.28E-125	AAEL010365-PA OS=Aedes aegypti GN=AAEL010365 PE=4 SV=1
CCG009637.1	825	397	2	20.6775571	0.084747338	-7.930681993	Down	1.34E-134	7.47E-134	AAEL013951-PA OS=Aedes aegypti GN=AAEL013951 PE=3 SV=1
CCG017474.1	363	437	2	51.729405	0.192607586	-8.069176266	Down	1.92E-148	1.09E-147	-
CCG018621.2	1665	445	2	11.4844035	0.041991924	-8.095348322	Down	3.27E-151	1.86E-150	-
CCG025854.1	1386	453	2	14.0442206	0.050444844	-8.121054036	Down	5.55E-154	3.17E-153	AAEL010365-PA OS=Aedes aegypti GN=AAEL010365 PE=4 SV=1
CCG019703.1	516	559	2	46.5505458	0.135497197	-8.424391269	Down	1.05E-190	6.25E-190	AAEL013449-PA OS=Aedes aegypti GN=AAEL013449 PE=3 SV=1

CCG007909.1	810	570	2	30.2379614	0.086316733	-8.452504905	Down	1.62E-194	9.63E-194	AAEL004659-PA OS=Aedes aegypti GN=AAEL004659 PE=4 SV=1
CCG027492.1	858	601	2	30.0988467	0.081487825	-8.528907977	Down	2.90E-205	1.74E-204	-
CCG006738.1	2055	608	2	12.7131867	0.034022654	-8.54561431	Down	1.09E-207	6.51E-207	AAEL003265-PA OS=Aedes aegypti GN=AAEL003265 PE=4 SV=1
CCG022601.1	633	1110	2	75.3497716	0.110452691	-9.414030757	Down	0	0	-
CCG012607.1	900	1135	2	54.1896098	0.077685059	-9.446163378	Down	0	0	AAEL009904-PA OS=Aedes aegypti GN=AAEL009904 PE=4 SV=1
CCG023020.1	177	1172	2	284.522763	0.395008777	-9.492443651	Down	0	0	AAEL006575-PA OS=Aedes aegypti GN=AAEL006575 PE=4 SV=1
CCG001594.1	744	1220	2	70.461124	0.093973862	-9.550352229	Down	0	0	AAEL009239-PA OS=Aedes aegypti GN=AAEL009239 PE=4 SV=1
CCG023070.1	906	1465	2	69.481966	0.077170589	-9.814371745	Down	0	0	-
CCG005289.1	468	1616	2	148.374126	0.149394345	-9.955898279	Down	0	0	AAEL002181-PA OS=Aedes aegypti GN=AAEL002181 PE=4 SV=1
CCG014525.1	624	1780	2	122.573922	0.112045759	-10.09534832	Down	0	0	AAEL010824-PA OS=Aedes aegypti GN=AAEL010824 PE=4 SV=1
CCG005201.1	471	2354	2	214.757442	0.148442789	-10.4985854	Down	0	0	AAEL006429-PA OS=Aedes aegypti GN=AAEL006429 PE=3 SV=1
CCG020273.1	381	2376	2	267.968739	0.183508015	-10.51200592	Down	0	0	Cuticle protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003476 PE=4 SV=1
CCG000986.1	564	5973	2	455.067775	0.12396552	-11.8419268	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010091 PE=4 SV=1
CCG005492.1	781	16	3	0.88030186	0.134282753	-2.712724295	Down	0.0005293	0.00083028	-
CCG002248.1	729	16	3	0.94309431	0.143861221	-2.712724295	Down	0.0005293	0.00083033	-
CCG015958.1	270	16	3	2.54635464	0.388425297	-2.712724295	Down	0.0005293	0.00083038	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009687 PE=4 SV=1
CCG013837.1	1752	16	3	0.39241767	0.059860063	-2.712724295	Down	0.0005293	0.00083049	AAEL000751-PA OS=Aedes aegypti GN=AAEL000751 PE=4 SV=1
CCG018818.1	615	16	3	1.11791179	0.170528179	-2.712724295	Down	0.0005293	0.00083044	AAEL013436-PA OS=Aedes aegypti GN=AAEL013436 PE=4 SV=1

CCG007990.1	690	17	3	1.05867462	0.151992508	-2.800187137	Down	0.000275	0.00043863	AGAP002453-PA OS=Anopheles gambiae GN=AgaP_AGAP002453 PE=4 SV=5
CCG028072.1	785	18	3	0.98529328	0.13359851	-2.882649297	Down	0.0001419	0.00022989	AAEL014449-PA OS=Aedes aegypti GN=AAEL014449 PE=4 SV=1
CCG015213.1	1071	19	3	0.76230155	0.097922344	-2.960651809	Down	7.28E-05	0.00012059	AAEL000268-PA OS=Aedes aegypti GN=AAEL000268 PE=3 SV=1
CCG022076.1	1314	19	3	0.62132797	0.079813417	-2.960651809	Down	7.28E-05	0.00012057	AAEL001168-PA OS=Aedes aegypti GN=AAEL001168 PE=4 SV=1
CCG012677.1	714	19	3	1.14345232	0.146883516	-2.960651809	Down	7.28E-05	0.00012058	-
CCG016764.1	708	20	3	1.21383431	0.148128291	-3.03465239	Down	3.71E-05	6.27E-05	AAEL015566-PA OS=Aedes aegypti GN=CTLGA2_b PE=4 SV=1
CCG005296.1	553	21	3	1.63176207	0.189647071	-3.105041718	Down	1.88E-05	3.24E-05	-
CCG002297.1	1941	21	3	0.46489667	0.05403134	-3.105041718	Down	1.88E-05	3.24E-05	AAEL002173-PA OS=Aedes aegypti GN=AAEL002173 PE=4 SV=1
CCG019946.1	827	22	3	1.14308847	0.12681358	-3.172155914	Down	9.50E-06	1.66E-05	Putative transposase OS=Anopheles gambiae PE=4 SV=1
CCG005286.1	315	22	3	3.00106083	0.332935969	-3.172155914	Down	9.50E-06	1.66E-05	AAEL011332-PA OS=Aedes aegypti GN=AAEL011332 PE=4 SV=1
CCG023443.1	1608	22	3	0.58789438	0.065220666	-3.172155914	Down	9.50E-06	1.66E-05	AAEL002666-PA OS=Aedes aegypti GN=AAEL002666 PE=4 SV=1
CCG014607.1	2994	23	3	0.33009482	0.035028333	-3.236286252	Down	4.78E-06	8.50E-06	AAEL009019-PA OS=Aedes aegypti GN=AAEL009019 PE=4 SV=1
CCG017407.1	1464	25	3	0.73377279	0.071635813	-3.356580485	Down	1.19E-06	2.18E-06	AAEL008715-PA OS=Aedes aegypti GN=AAEL008715 PE=4 SV=1
CCG009474.1	1137	25	3	0.94480507	0.092238197	-3.356580485	Down	1.19E-06	2.18E-06	Uncharacterized protein OS=Anopheles darlingi GN=AND_09382 PE=4 SV=1
CCG017624.1	1593	25	3	0.6743524	0.065834796	-3.356580485	Down	1.19E-06	2.18E-06	AAEL001312-PA OS=Aedes aegypti GN=CYP9M6 PE=3 SV=1
CCG022689.1	1398	26	3	0.799151	0.075017761	-3.413164014	Down	5.92E-07	1.10E-06	AAEL013156-PA OS=Aedes aegypti GN=AAEL013156 PE=4 SV=1

CCG017550.1	1539	26	3	0.72593444	0.068144789	-3.413164014	Down	5.92E-07	1.10E-06	MELZ16 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007168 PE=4 SV=1
CCG001215.1	1296	27	3	0.8952028	0.080921937	-3.467611798	Down	2.93E-07	5.51E-07	-
CCG001221.1	504	28	3	2.38720748	0.208084981	-3.520079218	Down	1.45E-07	2.75E-07	AAEL004969-PA OS=Aedes aegypti GN=AAEL004969 PE=4 SV=1
CCG020306.1	1392	28	3	0.86433374	0.075341114	-3.520079218	Down	1.45E-07	2.75E-07	AAEL002391-PA (Fragment) OS=Aedes aegypti GN=AAEL002391 PE=4 SV=1
CCG007459.1	309	29	3	4.03275826	0.339400745	-3.570705291	Down	7.12E-08	1.37E-07	-
CCG009504.1	1152	30	3	1.11900351	0.091037179	-3.619614891	Down	3.50E-08	6.83E-08	Lipase OS=Aedes aegypti GN=AAEL013365 PE=3 SV=1
CCG015428.1	2400	30	3	0.53712168	0.043697846	-3.619614891	Down	3.50E-08	6.83E-08	AAEL007730-PA OS=Aedes aegypti GN=AAEL007730 PE=4 SV=1
CCG011854.1	1008	31	3	1.32148985	0.10404249	-3.666920606	Down	1.71E-08	3.38E-08	-
CCG028569.1	843	31	3	1.58014445	0.124406679	-3.666920606	Down	1.71E-08	3.38E-08	-
CCG016193.1	840	34	3	1.73925116	0.124850988	-3.800187137	Down	1.98E-09	4.04E-09	AAEL011530-PA OS=Aedes aegypti GN=AAEL011530 PE=4 SV=1
CCG009995.1	669	34	3	2.18381312	0.156763573	-3.800187137	Down	1.98E-09	4.04E-09	AAEL012952-PA OS=Aedes aegypti GN=AAEL012952 PE=1 SV=1
CCG013221.1	1084	44	3	1.74415897	0.096747998	-4.172155914	Down	1.32E-12	3.20E-12	Copia OS=Drosophila simulans GN=GIP PE=4 SV=1
CCG014977.1	810	45	3	2.38720748	0.129475099	-4.204577392	Down	6.29E-13	1.60E-12	AAEL009485-PA OS=Aedes aegypti GN=AAEL009485 PE=4 SV=1
CCG018070.1	439	46	3	4.50252344	0.23889483	-4.236286252	Down	3.00E-13	8.29E-13	-
CCG014031.1	2211	47	3	0.91342267	0.047433211	-4.267313147	Down	1.43E-13	4.28E-13	AAEL002676-PA OS=Aedes aegypti GN=AAEL002676 PE=4 SV=1
CCG015690.1	1559	55	3	1.51593034	0.067270577	-4.494084009	Down	3.63E-16	1.36E-15	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007192 PE=4 SV=1
CCG020179.1	957	70	3	3.14303179	0.109587075	-4.842007312	Down	4.30E-21	1.70E-20	AAEL007582-PA OS=Aedes aegypti GN=AAEL007582 PE=3 SV=1
CCG025462.1	1059	76	3	3.0837581	0.099031946	-4.960651809	Down	4.43E-23	1.77E-22	AAEL007076-PA OS=Aedes aegypti GN=AAEL007076 PE=4 SV=1

CCG007094.1	1422	86	3	2.59873219	0.073751639	-5.13898905	Down	2.08E-26	8.58E-26	AAEL011374-PA OS=Aedes aegypti GN=AAEL011374 PE=4 SV=1
CCG000987.1	1608	90	3	2.40502246	0.065220666	-5.204577392	Down	9.62E-28	3.99E-27	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013304 PE=4 SV=1
CCG008093.1	582	114	3	8.41675214	0.180197303	-5.54561431	Down	8.40E-36	3.66E-35	Salivary antigen-5 related protein AG5-4 OS=Aedes albopictus PE=2 SV=1
CCG012197.1	907	117	3	5.54295364	0.115628258	-5.583089015	Down	8.18E-37	3.58E-36	AGAP013343-PA OS=Anopheles gambiae GN=AgaP_AGAP013343 PE=4 SV=1
CCG007229.1	1305	117	3	3.85245897	0.080363855	-5.583089015	Down	8.18E-37	3.58E-36	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018057 PE=4 SV=1
CCG011296.1	516	143	3	11.9082792	0.203245795	-5.872595632	Down	1.30E-45	5.96E-45	AGAP006830-PA OS=Anopheles gambiae GN=CPR58 PE=4 SV=3
CCG011984.1	1146	144	3	5.39933838	0.091513814	-5.882649297	Down	5.97E-46	2.74E-45	Proton channel protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001618 PE=4 SV=1
CCG019212.1	1770	149	3	3.61722625	0.059251317	-5.931892816	Down	1.20E-47	5.53E-47	AAEL011774-PA OS=Aedes aegypti GN=AAEL011774 PE=4 SV=1
CCG007391.1	621	175	3	12.1090234	0.168880564	-6.163935407	Down	1.70E-56	8.13E-56	-
CCG016297.1	3690	198	3	2.30569308	0.028421363	-6.342080916	Down	2.41E-64	1.17E-63	AAEL005967-PA OS=Aedes aegypti GN=AAEL005967 PE=4 SV=1
CCG016893.1	639	208	3	13.9870185	0.164123365	-6.413164014	Down	9.19E-68	4.53E-67	AGAP003937-PD OS=Anopheles gambiae GN=AgaP_AGAP003937 PE=4 SV=1
CCG023682.1	1290	215	3	7.16162244	0.081298318	-6.460917145	Down	3.70E-70	1.83E-69	AAEL006399-PA OS=Aedes aegypti GN=AAEL006399 PE=4 SV=1
CCG012992.1	747	283	3	16.2790293	0.140394686	-6.857382538	Down	1.77E-93	9.24E-93	Uncharacterized protein (Fragment) OS=Sarcophilus harrisii GN=PDIA4 PE=3 SV=1
CCG015822.1	621	320	3	22.1422143	0.168880564	-7.03465239	Down	3.34E-106	1.78E-105	-
CCG009040.1	486	374	3	33.0672443	0.215791832	-7.259618755	Down	8.42E-125	4.64E-124	AAEL002908-PA OS=Aedes aegypti GN=AAEL002908 PE=4 SV=1

CCG017754.1	576	530	3	39.5381239	0.182074358	-7.762572845	Down	1.16E-178	6.80E-178	AGAP013156-PA OS=Anopheles gambiae GN=AgaP_AGAP013156 PE=4 SV=1
CCG003582.1	2043	602	3	12.6616643	0.05133374	-7.946343972	Down	1.46E-203	8.72E-203	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018443 PE=4 SV=1
CCG023793.1	1881	622	3	14.2090244	0.055754827	-7.993495066	Down	1.75E-210	1.05E-209	AAEL003999-PA (Fragment) OS=Aedes aegypti GN=AAEL003999 PE=4 SV=1
CCG012638.1	549	1972	3	154.346661	0.191028835	-9.658168132	Down	0	0	AAEL013511-PA OS=Aedes aegypti GN=AAEL013511 PE=4 SV=1
CCG015981.1	1107	2518	3	97.7396493	0.094737877	-10.01078686	Down	0	0	AAEL003915-PA OS=Aedes aegypti GN=AAEL003915 PE=4 SV=1
CCG020824.1	504	4349	3	370.784476	0.208084981	-10.79919229	Down	0	0	-
CCG010041.1	543	5423	3	429.143408	0.193139651	-11.11759975	Down	0	0	AAEL013511-PA OS=Aedes aegypti GN=AAEL013511 PE=4 SV=1
CCG001372.1	609	5528	3	390.043831	0.17220826	-11.1452662	Down	0	0	AAEL004292-PA OS=Aedes aegypti GN=AAEL004292 PE=4 SV=1
CCG017458.1	597	12542	3	902.724307	0.175669733	-12.3272041	Down	0	0	AAEL010180-PA OS=Aedes aegypti GN=AAEL010180 PE=4 SV=1
CCG011223.1	430	18	4	1.79873308	0.325193272	-2.467611798	Down	0.0004703	0.00074043	-
CCG023256.1	775	18	4	0.99800674	0.180429816	-2.467611798	Down	0.0004703	0.00074029	-
CCG004106.1	1026	18	4	0.75385499	0.136289578	-2.467611798	Down	0.0004703	0.00074038	Lipase OS=Aedes aegypti GN=AAEL012350 PE=3 SV=1
CCG015861.1	534	18	4	1.44841802	0.261859751	-2.467611798	Down	0.0004703	0.00074034	AGAP003060-PA OS=Anopheles gambiae GN=AGAP003060 PE=4 SV=4
CCG028568.1	1152	19	4	0.70870222	0.121382905	-2.54561431	Down	0.0002511	0.00040181	AAEL002355-PA OS=Aedes aegypti GN=AAEL002355 PE=3 SV=1
CCG011894.1	1326	19	4	0.6157051	0.105454832	-2.54561431	Down	0.0002511	0.00040178	AAEL000347-PA OS=Aedes aegypti GN=AAEL000347 PE=4 SV=1
CCG009119.1	1736	19	4	0.47029087	0.080549025	-2.54561431	Down	0.0002511	0.00040183	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010720 PE=4 SV=1

CCG016042.1	894	19	4	0.91322702	0.156412871	-2.54561431	Down	0.0002511	0.00040176	AAEL005780-PA OS=Aedes aegypti GN=AAEL005780 PE=4 SV=1
CCG027667.1	605	20	4	1.42048709	0.231129103	-2.619614891	Down	0.0001331	0.00021597	Gag-Pol polyprotein OS=Schistosoma japonicum PE=4 SV=1
CCG017447.1	1713	20	4	0.50168984	0.081630535	-2.619614891	Down	0.0001331	0.000216	AAEL013428-PA OS=Aedes aegypti GN=AAEL013428 PE=4 SV=1
CCG016826.1	1884	20	4	0.4561543	0.074221394	-2.619614891	Down	0.0001331	0.00021598	AAEL008939-PA OS=Aedes aegypti GN=AAEL008939 PE=4 SV=1
CCG027183.2	1291	21	4	0.69896547	0.108313793	-2.690004219	Down	7.01E-05	0.00011624	DNA topoisomerase 1b OS=Megavirus chiliensis GN=mg859 PE=4 SV=1
CCG027926.1	315	21	4	2.86464897	0.443914626	-2.690004219	Down	7.01E-05	0.00011623	-
CCG025615.1	453	22	4	2.08683038	0.308682356	-2.757118415	Down	3.67E-05	6.20E-05	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010132 PE=4 SV=1
CCG000174.1	372	22	4	2.54122086	0.375895449	-2.757118415	Down	3.67E-05	6.20E-05	AAEL002191-PA OS=Aedes aegypti GN=AAEL002191 PE=4 SV=1
CCG022999.1	1014	22	4	0.93228221	0.137902472	-2.757118415	Down	3.67E-05	6.20E-05	AAEL002893-PA OS=Aedes aegypti GN=AAEL002893 PE=3 SV=1
CCG015509.1	880	23	4	1.12307261	0.158901258	-2.821248752	Down	1.91E-05	3.29E-05	-
CCG010727.1	861	23	4	1.14785586	0.16240779	-2.821248752	Down	1.91E-05	3.29E-05	AAEL012643-PA OS=Aedes aegypti GN=AAEL012643 PE=4 SV=1
CCG004851.1	540	25	4	1.98933957	0.258950198	-2.941542986	Down	5.09E-06	9.05E-06	AGAP013084-PA OS=Anopheles gambiae GN=AgaP_AGAP013084 PE=4 SV=1
CCG005600.1	1309	25	4	0.82065956	0.106824375	-2.941542986	Down	5.09E-06	9.05E-06	Pol-like protein OS=Aedes aegypti PE=4 SV=1
CCG006478.1	924	26	4	1.20910509	0.151334531	-2.998126514	Down	2.61E-06	4.70E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007764 PE=4 SV=1
CCG019890.1	1020	27	4	1.13743415	0.137091281	-3.052574298	Down	1.33E-06	2.44E-06	AAEL007386-PA OS=Aedes aegypti GN=AAEL007386 PE=3 SV=1
CCG006894.1	708	30	4	1.82075147	0.197504389	-3.204577392	Down	1.73E-07	3.29E-07	AAEL014235-PA OS=Aedes aegypti GN=AAEL014235 PE=4 SV=1

CCG019028.1	570	32	4	2.41233598	0.245321241	-3.297686796	Down	4.37E-08	8.52E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009310 PE=4 SV=1
CCG007576.1	2400	32	4	0.57292979	0.058263795	-3.297686796	Down	4.37E-08	8.52E-08	AAEL007730-PA OS=Aedes aegypti GN=AAEL007730 PE=4 SV=1
CCG003241.1	612	32	4	2.24678351	0.228485469	-3.297686796	Down	4.37E-08	8.52E-08	AAEL010851-PA OS=Aedes aegypti GN=AAEL010851 PE=4 SV=1
CCG016538.1	678	34	4	2.15482445	0.206243521	-3.385149637	Down	1.09E-08	2.17E-08	Lysozyme OS=Aedes albopictus PE=2 SV=1
CCG017216.1	2598	35	4	0.57888403	0.053823367	-3.426969813	Down	5.41E-09	1.09E-08	PTTBT OS=Culex quinquefasciatus GN=CpipJ_CPIJ007600 PE=4 SV=1
CCG010697.1	1149	36	4	1.34631022	0.121699832	-3.467611798	Down	2.69E-09	5.48E-09	Uncharacterized protein OS=Anopheles darlingi GN=AND_01397 PE=4 SV=1
CCG013645.1	804	36	4	1.92401797	0.173921775	-3.467611798	Down	2.69E-09	5.47E-09	AAEL009965-PA OS=Aedes aegypti GN=AAEL009965 PE=4 SV=1
CCG000520.1	864	36	4	1.79040561	0.161843874	-3.467611798	Down	2.69E-09	5.47E-09	AAEL008638-PA OS=Aedes aegypti GN=CYP49A1 PE=3 SV=1
CCG028016.1	1773	37	4	0.89671753	0.078868081	-3.507140162	Down	1.33E-09	2.73E-09	AAEL013187-PA (Fragment) OS=Aedes aegypti GN=AAEL013187 PE=4 SV=1
CCG016438.1	1068	48	4	1.93122403	0.130929876	-3.882649297	Down	5.05E-13	1.31E-12	AAEL004360-PA OS=Aedes aegypti GN=AAEL004360 PE=4 SV=1
CCG012696.1	771	48	4	2.67515858	0.181365898	-3.882649297	Down	5.05E-13	1.31E-12	AAEL009559-PA OS=Aedes aegypti GN=AAEL009559 PE=4 SV=1
CCG027851.1	453	50	4	4.74279631	0.308682356	-3.941542986	Down	1.18E-13	3.65E-13	AAEL000481-PA OS=Aedes aegypti GN=AAEL006481 PE=4 SV=1
CCG013449.1	1116	52	4	2.00217401	0.125298483	-3.998126514	Down	2.74E-14	9.56E-14	Hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG011893.1	780	53	4	2.91973838	0.179273214	-4.025607251	Down	1.32E-14	4.70E-14	AAEL000336-PA OS=Aedes aegypti GN=AAEL000336 PE=4 SV=1
CCG015219.1	924	58	4	2.69723442	0.151334531	-4.155667791	Down	3.35E-16	1.26E-15	AAEL000098-PA OS=Aedes aegypti GN=AAEL000098 PE=4 SV=1

CCG023314.1	1305	73	4	2.40367098	0.107151806	-4.487511355	Down	4.76E-21	1.87E-20	AAEL014228-PA (Fragment) OS=Aedes aegypti GN=AAEL014228 PE=3 SV=1
CCG020354.1	573	82	4	6.14924649	0.244036836	-4.655238801	Down	5.44E-24	2.19E-23	Uncharacterized protein OS=Anopheles darlingi GN=AND_08994 PE=4 SV=1
CCG025404.1	988	86	4	3.74028054	0.141531485	-4.723951551	Down	2.64E-25	1.08E-24	-
CCG015540.1	651	91	4	6.00652204	0.2147974	-4.805481436	Down	5.95E-27	2.46E-26	AAEL010545-PA (Fragment) OS=Aedes aegypti GN=AAEL010545 PE=4 SV=1
CCG021843.1	410	97	4	10.1660104	0.341056359	-4.897599638	Down	6.19E-29	2.59E-28	AAEL000578-PA OS=Aedes aegypti GN=AAEL000578 PE=4 SV=1
CCG001422.1	855	99	4	4.97544295	0.163547494	-4.927043416	Down	1.35E-29	5.66E-29	AAEL008421-PA OS=Aedes aegypti GN=AAEL008421 PE=4 SV=1
CCG017063.1	573	105	4	7.87403514	0.244036836	-5.011932314	Down	1.38E-31	5.85E-31	Uncharacterized protein OS=Anopheles darlingi GN=AND_08994 PE=4 SV=1
CCG016445.1	1434	112	4	3.35607411	0.097512627	-5.105041718	Down	6.44E-34	2.77E-33	AAEL005004-PA OS=Aedes aegypti GN=AAEL005004 PE=4 SV=1
CCG027136.1	960	124	4	5.55025739	0.145659487	-5.251883107	Down	6.34E-38	2.80E-37	AAEL009697-PA OS=Aedes aegypti GN=AAEL009697 PE=4 SV=1
CCG008192.1	1299	127	4	4.20104411	0.107646734	-5.286371483	Down	6.28E-39	2.79E-38	AAEL004270-PA OS=Aedes aegypti GN=AAEL004270 PE=4 SV=1
CCG013835.1	1719	128	4	3.19960793	0.081345612	-5.297686796	Down	2.90E-39	1.29E-38	AAEL002405-PA OS=Aedes aegypti GN=AAEL002405 PE=4 SV=1
CCG018885.1	930	159	4	7.3464385	0.15035818	-5.610569751	Down	1.09E-49	5.06E-49	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002954 PE=4 SV=1
CCG020181.1	504	176	4	15.0053041	0.277446641	-5.757118415	Down	1.95E-55	9.26E-55	AAEL011591-PA OS=Aedes aegypti GN=AAEL011591 PE=4 SV=1
CCG005809.1	1554	180	4	4.97718934	0.089982694	-5.789539893	Down	8.61E-57	4.11E-56	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013980 PE=4 SV=1

CCG021901.1	576	213	4	15.8898498	0.242765811	-6.032396416	Down	5.40E-68	2.66E-67	AGAP013156-PA OS=Anopheles gambiae GN=AgaP_AGAP013156 PE=4 SV=1
CCG002556.1	1254	244	4	8.3609372	0.111509655	-6.228424134	Down	1.48E-78	7.51E-78	AAEL007887-PA OS=Aedes aegypti GN=AAEL007887 PE=3 SV=1
CCG009157.1	1218	262	4	9.2430792	0.114805507	-6.331109798	Down	1.06E-84	5.45E-84	Lipase OS=Aedes aegypti GN=AAEL012342 PE=3 SV=1
CCG015682.1	933	303	4	13.9548013	0.149874713	-6.54086078	Down	1.00E-98	5.30E-98	AAEL000694-PA OS=Aedes aegypti GN=AAEL000694 PE=4 SV=1
CCG011371.1	297	370	4	53.5313192	0.470818542	-6.829068257	Down	1.04E-121	5.72E-121	AAEL006011-PA OS=Aedes aegypti GN=AAEL006011 PE=4 SV=1
CCG015691.1	1881	418	4	9.54882991	0.07433977	-7.005045928	Down	3.29E-138	1.85E-137	Uncharacterized protein OS=Anopheles darlingi GN=AND_13394 PE=4 SV=1
CCG024842.1	651	508	4	33.5309143	0.2147974	-7.286371483	Down	3.31E-169	1.92E-168	Uncharacterized protein OS=Anopheles darlingi GN=AND_08375 PE=4 SV=1
CCG015458.1	540	560	4	44.5612063	0.258950198	-7.426969813	Down	3.84E-187	2.27E-186	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001222 PE=4 SV=1
CCG013055.1	486	657	4	58.0887153	0.287722443	-7.657436357	Down	1.23E-220	7.43E-220	AAEL002908-PA OS=Aedes aegypti GN=AAEL002908 PE=4 SV=1
CCG025647.1	591	800	4	58.1654614	0.236604242	-7.941542986	Down	4.42E-270	2.75E-269	AAEL009707-PA OS=Aedes aegypti GN=AAEL009707 PE=4 SV=1
CCG023324.1	1388	1151	4	35.6326834	0.100744313	-8.466358914	Down	0	0	AAEL005188-PA OS=Aedes aegypti GN=AAEL005188 PE=4 SV=1
CCG009378.1	484	1837	4	163.089675	0.288911378	-9.140822707	Down	0	0	AAEL005219-PA OS=Aedes aegypti GN=AAEL005219 PE=4 SV=1
CCG007385.1	222	2061	4	398.921725	0.629878861	-9.306815586	Down	0	0	AAEL006746-PA OS=Aedes aegypti GN=AAEL006746 PE=4 SV=1
CCG001090.1	3984	4927	4	53.1405327	0.035098671	-10.56418055	Down	0	0	AAEL000637-PA OS=Aedes aegypti GN=AAEL000637 PE=4 SV=1
CCG015425.1	1377	21	5	0.65531186	0.126936372	-2.368076124	Down	0.0002171	0.00034822	AAEL001689-PA OS=Aedes aegypti GN=AAEL001689 PE=3 SV=1

CCG006507.1	884	21	5	1.02077424	0.19772781	-2.368076124	Down	0.0002171	0.00034828	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG010239.1	895	21	5	1.00822841	0.195297636	-2.368076124	Down	0.0002171	0.00034824	-
CCG000127.1	2067	21	5	0.43655754	0.084562837	-2.368076124	Down	0.0002171	0.00034826	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002423 PE=4 SV=1
CCG020542.1	198	22	5	4.77441496	0.882784767	-2.43519032	Down	0.0001176	0.0001913	AGAP006541-PA OS=Anopheles gambiae GN=AGAP006541 PE=4 SV=3
CCG026081.1	966	22	5	0.97860679	0.180943462	-2.43519032	Down	0.0001176	0.00019131	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016285 PE=4 SV=1
CCG000494.1	1061	23	5	0.93148341	0.164742115	-2.499320657	Down	6.33E-05	0.00010519	Pol polyprotein (Fragment) OS=Anopheles gambiae PE=4 SV=1
CCG017427.1	424	23	5	2.33090542	0.41224383	-2.499320657	Down	6.33E-05	0.00010518	-
CCG027205.1	1314	23	5	0.75213386	0.133022362	-2.499320657	Down	6.33E-05	0.00010518	AAEL000048-PA (Fragment) OS=Aedes aegypti GN=GPRgr4 PE=4 SV=2
CCG015659.1	591	27	5	1.96308432	0.295755303	-2.730646203	Down	5.00E-06	8.89E-06	AAEL003184-PB OS=Aedes aegypti GN=AAEL003184 PE=4 SV=1
CCG026077.1	753	27	5	1.54074746	0.232126672	-2.730646203	Down	5.00E-06	8.89E-06	AGAP000801-PB (Fragment) OS=Anopheles gambiae GN=AgaP_AGAP000801 PE=4 SV=1
CCG019456.1	1875	27	5	0.61876418	0.093222071	-2.730646203	Down	5.00E-06	8.89E-06	AAEL013428-PA OS=Aedes aegypti GN=AAEL013428 PE=4 SV=1
CCG011905.1	624	27	5	1.85926736	0.280114397	-2.730646203	Down	5.00E-06	8.89E-06	-
CCG027282.1	876	33	5	1.61872288	0.199533543	-3.020152821	Down	9.63E-08	1.85E-07	AAEL007878-PA OS=Aedes aegypti GN=AAEL007878 PE=3 SV=1
CCG013797.1	756	33	5	1.87566302	0.231205534	-3.020152821	Down	9.63E-08	1.85E-07	-
CCG005675.1	1087	34	5	1.34403954	0.160801641	-3.063221543	Down	4.92E-08	9.57E-08	-
CCG003204.1	1638	35	5	0.91815672	0.106710247	-3.105041718	Down	2.51E-08	4.93E-08	AAEL005060-PA OS=Aedes aegypti GN=CLIPB44 PE=3 SV=2

CCG004659.1	564	38	5	2.89512396	0.309913801	-3.223686215	Down	3.25E-09	6.61E-09	AAEL001552-PA OS=Aedes aegypti GN=AAEL001552 PE=4 SV=1
CCG007486.3	1113	39	5	1.50567803	0.157045269	-3.26116092	Down	1.64E-09	3.35E-09	AAEL009774-PA OS=Aedes aegypti GN=AAEL009774 PE=3 SV=1
CCG005553.1	234	40	5	7.34525378	0.746971726	-3.297686796	Down	8.22E-10	1.70E-09	AAEL0005257-PA OS=Aedes aegypti GN=AAEL003257 PE=4 SV=1
CCG011864.1	984	41	5	1.79040561	0.17763352	-3.333310706	Down	4.11E-10	8.58E-10	AAEL011365-PA OS=Aedes aegypti GN=AAEL011365 PE=4 SV=1
CCG022553.1	436	52	5	5.12483073	0.400897669	-3.676198419	Down	1.78E-13	5.20E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018057 PE=4 SV=1
CCG007060.1	3015	62	5	0.88362307	0.057973925	-3.929955012	Down	1.32E-16	4.98E-16	AAEL0006541-PA OS=Aedes aegypti GN=AAEL006541 PE=4 SV=1
CCG019697.2	1170	63	5	2.31375494	0.149394345	-3.953038625	Down	6.39E-17	2.41E-16	AAEL0006826-PA OS=Aedes aegypti GN=AAEL006826 PE=3 SV=1
CCG018754.1	2073	80	5	1.65826279	0.084318082	-4.297686796	Down	2.36E-22	9.38E-22	Flu-1esmn OS=Culex quinquefasciatus GN=CpipJ_CPIJ001605 PE=4 SV=1
CCG014853.1	1143	82	5	3.08269312	0.152923345	-4.333310706	Down	5.33E-23	2.13E-22	AAEL014193-PA OS=Aedes aegypti GN=AAEL014193 PE=4 SV=1
CCG009581.1	528	93	5	7.5685328	0.331044288	-4.514917512	Down	1.43E-26	5.90E-26	AAEL013621-PA OS=Aedes aegypti GN=AAEL013621 PE=4 SV=1
CCG006055.1	738	114	5	6.63760128	0.236844694	-4.808648715	Down	1.85E-33	7.94E-33	AAEL010397-PA OS=Aedes aegypti GN=AAEL010397 PE=4 SV=1
CCG013201.1	1170	122	5	4.48060481	0.149394345	-4.906496039	Down	4.20E-36	1.83E-35	AAEL006154-PA (Fragment) OS=Aedes aegypti GN=AAEL006154 PE=4 SV=1
CCG018004.1	3975	122	5	1.31881953	0.043972675	-4.906496039	Down	4.20E-36	1.83E-35	AAEL015231-PA OS=Aedes aegypti GN=AAEL015231 PE=4 SV=1
CCG014391.1	2352	144	5	2.63080008	0.074316065	-5.145683703	Down	2.05E-43	9.28E-43	AAEL0006541-PA OS=Aedes aegypti GN=AAEL006541 PE=4 SV=1
CCG025432.1	1728	157	5	3.9040789	0.101152421	-5.27037945	Down	9.28E-48	4.29E-47	AAEL001347-PA OS=Aedes aegypti GN=AAEL001347 PE=4 SV=1

CCG022313.1	2727	176	5	2.77325753	0.064096584	-5.43519032	Down	3.93E-54	1.86E-53	AAEL001340-PA (Fragment) OS=Aedes aegypti GN=AAEL001340 PE=4 SV=1
CCG010418.1	1326	304	5	9.85128154	0.13181854	-6.223686215	Down	1.56E-97	8.24E-97	AAEL004939-PA OS=Aedes aegypti GN=AAEL004939 PE=3 SV=1
CCG008346.1	13800	1000	5	3.11374888	0.012666042	-7.941542986	Down	0	0	AAEL007225-PA OS=Aedes aegypti GN=AAEL007225 PE=4 SV=1
CCG024336.1	2091	1064	5	21.8650395	0.083592245	-8.031041137	Down	0	0	AAEL014174-PA OS=Aedes aegypti GN=AAEL014174 PE=4 SV=1
CCG009693.1	576	1384	5	103.246723	0.303457264	-8.410386929	Down	0	0	AAEL001677-PA OS=Aedes aegypti GN=AAEL001677 PE=4 SV=1
CCG003394.1	1155	2180	5	81.1030489	0.151334531	-9.065871121	Down	0	0	AAEL003915-PA OS=Aedes aegypti GN=AAEL003915 PE=4 SV=1
CCG007995.1	1623	6100	5	161.500543	0.107696478	-10.55035223	Down	0	0	Cytosol aminopeptidase OS=Culex quinquefasciatus GN=CpipJ_CPIJ000990 PE=4 SV=1
CCG024515.1	1027	21	6	0.87864112	0.204235307	-2.105041718	Down	0.0005818	0.00091085	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG008140.1	779	22	6	1.21352267	0.26925502	-2.172155914	Down	0.0003259	0.00051726	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG022683.1	660	23	6	1.49743015	0.317802516	-2.236286252	Down	0.0001811	0.0002917	AAEL008654-PA OS=Aedes aegypti GN=AAEL008654 PE=4 SV=1
CCG012465.1	618	23	6	1.59919724	0.339400745	-2.236286252	Down	0.0001811	0.00029172	Ecdysis triggering hormone OS=Aedes aegypti PE=2 SV=1
CCG000151.1	1068	24	6	0.96561201	0.196394813	-2.297686796	Down	9.99E-05	0.00016297	AAEL007851-PA OS=Aedes aegypti GN=AAEL007851 PE=4 SV=1
CCG005775.1	1818	24	6	0.56725722	0.115373851	-2.297686796	Down	9.99E-05	0.00016296	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC011569 PE=4 SV=1
CCG013605.1	321	24	6	3.21269044	0.653425734	-2.297686796	Down	9.99E-05	0.00016295	AAEL012977-PA (Fragment) OS=Aedes aegypti GN=AAEL012977 PE=4 SV=1
CCG026627.1	436	25	6	2.46386093	0.481077203	-2.356580485	Down	5.47E-05	9.13E-05	-

CCG021796.1	600	25	6	1.79040561	0.349582768	-2.356580485	Down	5.47E-05	9.13E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011650 PE=4 SV=1
CCG003705.1	896	26	6	1.24688962	0.234095603	-2.413164014	Down	2.98E-05	5.06E-05	-
CCG008952.1	1404	26	6	0.79573583	0.149394345	-2.413164014	Down	2.98E-05	5.06E-05	AAEL010193-PA OS=Aedes aegypti GN=AAEL010193 PE=4 SV=1
CCG004928.1	747	26	6	1.49559987	0.280789372	-2.413164014	Down	2.98E-05	5.06E-05	AAEL013737-PA OS=Aedes aegypti GN=AAEL013737 PE=4 SV=1
CCG015038.1	481	27	6	2.41202252	0.436069981	-2.467611798	Down	1.61E-05	2.78E-05	AAEL010836-PA OS=Aedes aegypti GN=AAEL010836 PE=4 SV=1
CCG024862.1	543	30	6	2.37401849	0.386279301	-2.619614891	Down	2.47E-06	4.46E-06	AAEL000557-PA OS=Aedes aegypti GN=AAEL000557 PE=4 SV=1
CCG011169.1	1185	30	6	1.08784138	0.177003933	-2.619614891	Down	2.47E-06	4.46E-06	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004277 PE=4 SV=1
CCG005125.1	3588	30	6	0.35927872	0.058458657	-2.619614891	Down	2.47E-06	4.46E-06	Carboxypeptidase AT OS=Culex quinquefasciatus GN=CpipJ_CPIJ007169 PE=3 SV=1
CCG020602.1	465	31	6	2.86464897	0.451074539	-2.666920606	Down	1.31E-06	2.40E-06	AAEL002438-PA (Fragment) OS=Aedes aegypti GN=AAEL002438 PE=4 SV=1
CCG006172.1	537	31	6	2.48056196	0.390595271	-2.666920606	Down	1.31E-06	2.40E-06	AAEL011373-PA OS=Aedes aegypti GN=AAEL011373 PE=4 SV=1
CCG003569.1	588	32	6	2.33848896	0.35671711	-2.712724295	Down	6.92E-07	1.28E-06	AAEL001305-PA OS=Aedes aegypti GN=AAEL001305 PE=4 SV=1
CCG001953.1	960	32	6	1.43232449	0.21848923	-2.712724295	Down	6.92E-07	1.28E-06	-
CCG020517.1	729	34	6	2.00407541	0.287722443	-2.800187137	Down	1.90E-07	3.60E-07	AAEL014538-PA OS=Aedes aegypti GN=AAEL014538 PE=4 SV=1
CCG005252.1	726	36	6	2.13073064	0.288911378	-2.882649297	Down	5.15E-08	1.00E-07	AAEL012748-PA OS=Aedes aegypti GN=AAEL012748 PE=4 SV=1
CCG013483.1	1029	37	6	1.54507306	0.203838349	-2.922177661	Down	2.66E-08	5.24E-08	AAEL009690-PA OS=Aedes aegypti GN=AAEL009690 PE=4 SV=1

CCG005045.1	747	38	6	2.18587673	0.280789372	-2.960651809	Down	1.37E-08	2.73E-08	Uncharacterized protein OS=Anopheles darlingi GN=AND_17075 PE=4 SV=1
CCG022710.1	1485	38	6	1.09956223	0.141245563	-2.960651809	Down	1.37E-08	2.73E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008816 PE=4 SV=1
CCG024771.1	603	43	6	3.06417676	0.34784355	-3.13898905	Down	4.80E-10	1.00E-09	-
CCG006183.1	558	47	6	3.61931456	0.375895449	-3.267313147	Down	3.13E-11	6.80E-11	-
CCG023284.1	621	48	6	3.32133214	0.337761128	-3.297686796	Down	1.57E-11	3.47E-11	AAEL006974-PA OS=Aedes aegypti GN=AAEL006974 PE=3 SV=1
CCG020526.1	726	52	6	3.07772204	0.288911378	-3.413164014	Down	9.83E-13	2.43E-12	AAEL006310-PA OS=Aedes aegypti GN=AAEL006310 PE=4 SV=1
CCG020296.1	363	54	6	6.39219193	0.577822757	-3.467611798	Down	2.43E-13	6.84E-13	AAEL017144-PA OS=Aedes aegypti PE=4 SV=1
CCG027248.1	345	55	6	6.85024755	0.607970031	-3.494084009	Down	1.20E-13	3.71E-13	-
CCG015057.1	729	57	6	3.35977349	0.287722443	-3.54561431	Down	2.95E-14	1.02E-13	AAEL013773-PA OS=Aedes aegypti GN=AAEL013773 PE=4 SV=1
CCG014035.1	2055	59	6	1.23368095	0.102067961	-3.595367345	Down	7.17E-15	2.57E-14	AAEL013187-PA (Fragment) OS=Aedes aegypti GN=AAEL013187 PE=4 SV=1
CCG010071.1	1125	70	6	2.67367238	0.186444143	-3.842007312	Down	2.73E-18	1.05E-17	AAEL006484-PA OS=Aedes aegypti GN=GPROP3 PE=3 SV=1
CCG018499.1	957	78	6	3.50223542	0.219174149	-3.998126514	Down	8.21E-21	3.23E-20	AAEL014757-PA OS=Aedes aegypti GN=AAEL014757 PE=4 SV=1
CCG013843.1	384	80	6	8.95202804	0.546223075	-4.03465239	Down	1.90E-21	7.52E-21	AAEL002022-PA OS=Aedes aegypti GN=AAEL002022 PE=4 SV=1
CCG017702.1	1938	90	6	1.99549851	0.108229959	-4.204577392	Down	1.22E-24	4.95E-24	AAEL006732-PA OS=Aedes aegypti GN=AAEL006732 PE=4 SV=1
CCG002127.1	588	94	6	6.86931132	0.35671711	-4.267313147	Down	6.33E-26	2.59E-25	-
CCG025477.1	1071	118	6	4.73429382	0.195844688	-4.595367345	Down	1.02E-33	4.39E-33	AAEL002911-PA (Fragment) OS=Aedes aegypti GN=AAEL002911 PE=4 SV=1
CCG022856.1	1875	146	6	3.34591	0.111866486	-4.902548854	Down	6.24E-43	2.82E-42	heat shock protein 70 BZ OS=Culex quinquefasciatus GN=CpipJ_CPIJ014564 PE=3 SV=1

CCG014629.1	615	148	6	10.3406841	0.341056359	-4.922177661	Down	1.36E-43	6.15E-43	AAEL004261-PA OS=Aedes aegypti GN=AAEL004261 PE=4 SV=1
CCG002211.1	568	156	6	11.8015468	0.369277572	-4.998126514	Down	3.02E-46	1.39E-45	-
CCG021128.2	381	157	6	17.7066885	0.550524044	-5.007345044	Down	1.41E-46	6.47E-46	AAEL001963-PA OS=Aedes aegypti GN=AAEL001963 PE=4 SV=1
CCG009629.1	1545	158	6	4.3943159	0.135760298	-5.016505044	Down	6.55E-47	3.01E-46	AAEL004564-PA (Fragment) OS=Aedes aegypti GN=AAEL004564 PE=4 SV=1
CCG023935.1	2064	160	6	3.33098718	0.101622898	-5.03465239	Down	1.42E-47	6.54E-47	AAEL010044-PA OS=Aedes aegypti GN=AAEL010044 PE=4 SV=1
CCG009570.1	429	165	6	16.526821	0.488926948	-5.07904651	Down	3.08E-49	1.43E-48	-
CCG020744.1	768	170	6	9.5115298	0.273111537	-5.122115232	Down	6.65E-51	3.11E-50	AAEL014840-PA OS=Aedes aegypti GN=AAEL014840 PE=3 SV=1
CCG001053.1	1077	170	6	6.78259506	0.194753631	-5.122115232	Down	6.65E-51	3.11E-50	AAEL009514-PA OS=Aedes aegypti GN=AAEL009514 PE=4 SV=1
CCG028421.1	1272	174	6	5.87793539	0.164897532	-5.155667791	Down	3.08E-52	1.45E-51	AAEL001437-PA OS=Aedes aegypti GN=AAEL001437 PE=4 SV=1
CCG015289.1	1227	206	6	7.21415267	0.170945119	-5.399224823	Down	5.98E-63	2.91E-62	AAEL012424-PA OS=Aedes aegypti GN=AAEL012424 PE=3 SV=1
CCG009243.1	1041	220	6	9.0810198	0.201488627	-5.494084009	Down	1.17E-67	5.78E-67	AAEL012424-PA OS=Aedes aegypti GN=AAEL012424 PE=3 SV=1
CCG010809.1	1395	234	6	7.20782645	0.15035818	-5.583089015	Down	2.25E-72	1.12E-71	AAEL009236-PA OS=Aedes aegypti GN=AAEL009236 PE=3 SV=1
CCG009382.1	330	247	6	32.1621953	0.635605032	-5.661091527	Down	9.22E-77	4.65E-76	AAEL011558-PA OS=Aedes aegypti GN=AAEL011558 PE=3 SV=1
CCG000684.1	489	297	6	26.0981824	0.428935911	-5.927043416	Down	1.07E-93	5.59E-93	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC005836 PE=4 SV=1
CCG006979.1	618	405	6	28.1597775	0.339400745	-6.374502393	Down	1.68E-130	9.33E-130	-
CCG027117.1	2142	414	6	8.30507476	0.097922344	-6.406211253	Down	1.41E-133	7.85E-133	AAEL010682-PA OS=Aedes aegypti GN=AAEL010682 PE=4 SV=1
CCG016902.1	5142	517	6	4.32037199	0.040791455	-6.726744766	Down	7.27E-169	4.22E-168	AAEL005109-PA OS=Aedes aegypti GN=AAEL005109 PE=4 SV=1

CCG018255.1	1890	756	6	17.1878938	0.110978656	-7.27496672	Down	4.34E-251	2.67E-250	AAEL009204-PA OS=Aedes aegypti GN=AAEL009204 PE=3 SV=1
CCG014812.1	222	887	6	171.685381	0.944818291	-7.50551459	Down	2.78E-296	1.75E-295	AAEL006746-PA OS=Aedes aegypti GN=AAEL006746 PE=4 SV=1
CCG005200.1	759	2170	6	122.851547	0.276350014	-8.796203623	Down	0	0	AAEL006403-PA OS=Aedes aegypti GN=AAEL006403 PE=3 SV=1
CCG020271.1	387	2738	6	304.008097	0.541988787	-9.131631027	Down	0	0	AAEL002181-PA OS=Aedes aegypti GN=AAEL002181 PE=4 SV=1
CCG002696.1	909	3027	6	143.090634	0.230747701	-9.276397255	Down	0	0	AAEL006179-PA (Fragment) OS=Aedes aegypti GN=AAEL006179 PE=3 SV=1
CCG020535.1	927	4796	6	222.311594	0.226267164	-9.940340239	Down	0	0	AAEL003027-PA OS=Aedes aegypti GN=AAEL003027 PE=4 SV=1
CCG011577.1	1017	24	7	1.01403504	0.240617441	-2.075294375	Down	0.0002612	0.00041756	AAEL013337-PA OS=Aedes aegypti GN=AAEL013337 PE=4 SV=1
CCG022894.1	1041	24	7	0.99065671	0.235070065	-2.075294375	Down	0.0002612	0.00041753	AAEL009072-PA OS=Aedes aegypti GN=AAEL009072 PE=4 SV=1
CCG010355.1	1545	24	7	0.66749102	0.158387015	-2.075294375	Down	0.0002612	0.00041759	AAEL014020-PA OS=Aedes aegypti GN=AAEL014020 PE=4 SV=1
CCG007276.1	846	25	7	1.26979121	0.289252881	-2.134188064	Down	0.0001473	0.00023834	AAEL008668-PA OS=Aedes aegypti GN=CLIPB22 PE=3 SV=1
CCG011644.1	336	27	7	3.4529251	0.728297433	-2.245219376	Down	4.59E-05	7.72E-05	AAEL000266-PA (Fragment) OS=Aedes aegypti GN=GPRNNA21 PE=4 SV=1
CCG015597.1	396	27	7	2.92975463	0.617949337	-2.245219376	Down	4.59E-05	7.72E-05	-
CCG007346.1	909	28	7	1.32360019	0.269205652	-2.297686796	Down	2.54E-05	4.35E-05	AAEL005624-PA OS=Aedes aegypti GN=AAEL005624 PE=4 SV=1
CCG027964.1	3357	28	7	0.35840112	0.072894828	-2.297686796	Down	2.54E-05	4.35E-05	AAEL007276-PA OS=Aedes aegypti GN=AAEL007276 PE=4 SV=1
CCG017478.1	663	29	7	1.87952082	0.369091912	-2.348312869	Down	1.39E-05	2.42E-05	Superoxide dismutase [Cu-Zn] OS=Aedes aegypti GN=CUSOD1 PE=3 SV=2
CCG017112.1	1602	33	7	0.88514435	0.152751521	-2.534725993	Down	1.20E-06	2.21E-06	AAEL000508-PA OS=Aedes aegypti GN=AAEL000508 PE=4 SV=1

CCG019465.1	1242	34	7	1.17630513	0.197027325	-2.577794715	Down	6.45E-07	1.20E-06	AAEL008558-PA (Fragment) OS=Aedes aegypti GN=AAEL008558 PE=4 SV=1
CCG028257.1	873	36	7	1.77194782	0.280306916	-2.660256876	Down	1.83E-07	3.47E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013073 PE=4 SV=1
CCG018394.1	435	36	7	3.55611597	0.562546983	-2.660256876	Down	1.83E-07	3.47E-07	Serine/threonine-protein phosphatase OS=Anopheles darlingi GN=AND_16101 PE=3 SV=1
CCG020162.1	1199	37	7	1.32600515	0.204093359	-2.69978524	Down	9.67E-08	1.86E-07	Retrovirus-related Pol polyprotein from transposon TNT 1-94 OS=Trichinella spiralis GN=Tsp_01017 PE=4 SV=1
CCG009470.1	1789	42	7	1.00879198	0.136784761	-2.882649297	Down	3.80E-09	7.68E-09	AAEL008340-PA OS=Aedes aegypti GN=AAEL008346 PE=4 SV=1
CCG019171.1	1191	45	7	1.62354161	0.205464263	-2.98218497	Down	5.24E-10	1.09E-09	AAEL003719-PA OS=Aedes aegypti GN=AAEL003719 PE=4 SV=1
CCG015306.1	657	47	7	3.0739384	0.372462614	-3.044920726	Down	1.38E-10	2.94E-10	AAEL011897-PA OS=Aedes aegypti GN=AAEL011897 PE=4 SV=1
CCG011507.1	5337	51	7	0.41061579	0.045851216	-3.162757216	Down	9.32E-12	2.08E-11	AAEL007856-PA OS=Aedes aegypti GN=AAEL007856 PE=4 SV=1
CCG013558.1	372	53	7	6.12203208	0.657817036	-3.218252329	Down	2.39E-12	5.64E-12	AAEL002191-PA OS=Aedes aegypti GN=AAEL002191 PE=4 SV=1
CCG015775.1	348	57	7	7.03814619	0.703183728	-3.323221888	Down	1.53E-13	4.55E-13	-
CCG015776.1	838	58	7	2.97403891	0.292014245	-3.348312869	Down	7.67E-14	2.47E-13	-
CCG006902.1	918	64	7	2.99571135	0.266566381	-3.490331874	Down	1.17E-15	4.26E-15	AAEL004270-PA OS=Aedes aegypti GN=AAEL004270 PE=4 SV=1
CCG007787.1	456	66	7	6.21930369	0.536640214	-3.534725993	Down	2.87E-16	1.08E-15	AAEL011730-PA OS=Aedes aegypti GN=AAEL011730 PE=4 SV=1
CCG005406.1	2263	72	7	1.36713252	0.108134307	-3.660256876	Down	4.10E-18	1.57E-17	AAEL014339-PA OS=Aedes aegypti GN=AAEL014339 PE=4 SV=1
CCG017995.1	780	73	7	4.02152644	0.313728125	-3.680156433	Down	2.01E-18	7.72E-18	AAEL003030-PA OS=Aedes aegypti GN=AAEL003030 PE=4 SV=1

CCG009819.1	1428	80	7	2.40726805	0.171364102	-3.812259969	Down	1.33E-20	5.24E-20	Uncharacterized protein OS=Anopheles darlingi GN=AND_13756 PE=4 SV=1
CCG015897.1	1389	92	7	2.84608753	0.176175621	-4.01389383	Down	2.21E-24	8.95E-24	AAEL006700-PA OS=Aedes aegypti GN=AAEL006700 PE=3 SV=1
CCG002751.1	792	153	7	8.30097146	0.308974668	-4.747719717	Down	3.81E-44	1.73E-43	AAEL002096-PA OS=Aedes aegypti GN=AAEL002096 PE=4 SV=1
CCG000271.1	1893	160	7	3.63188459	0.129269909	-4.812259969	Down	1.89E-46	8.66E-46	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008823 PE=4 SV=1
CCG027989.1	795	246	7	13.2962952	0.307808726	-5.432846379	Down	4.03E-75	2.02E-74	AAEL006421-PA OS=Aedes aegypti GN=AAEL006421 PE=3 SV=1
CCG013532.1	3114	278	7	3.83609063	0.078583153	-5.609272947	Down	6.73E-86	3.46E-85	Uncharacterized protein OS=Anopheles darlingi GN=AND_00371 PE=4 SV=1
CCG009811.1	1182	533	7	19.3763693	0.207028712	-6.548323597	Down	1.00E-172	5.82E-172	AAEL007126-PA OS=Aedes aegypti GN=AAEL007126 PE=4 SV=1
CCG009055.1	1686	793	7	20.2105573	0.145141125	-7.12150893	Down	4.79E-262	2.96E-261	AAEL009910-PA OS=Aedes aegypti GN=AAEL009910 PE=4 SV=1
CCG022356.1	696	1836	7	113.351196	0.351591864	-8.332682218	Down	0	0	AAEL003501-PA OS=Aedes aegypti GN=AAEL003501 PE=4 SV=1
CCG023071.1	933	8725	7	401.833799	0.262280747	-10.58127129	Down	0	0	-
CCG012823.1	1131	31923	7	1212.84071	0.216364224	-12.45264049	Down	0	0	AAEL009451-PA OS=Aedes aegypti GN=AAEL009451 PE=3 SV=1
CCG006452.1	816	24	8	1.26381572	0.342728204	-1.882649297	Down	0.0006169	0.00095923	Uncharacterized protein OS=Anopheles darlingi GN=AND_06563 PE=3 SV=1
CCG016083.1	369	24	8	2.79477949	0.757903019	-1.882649297	Down	0.0006169	0.00095917	-
CCG028646.1	357	25	8	3.00908506	0.783378751	-1.941542986	Down	0.000358	0.00056684	-
CCG001065.1	1596	25	8	0.67308482	0.175229458	-1.941542986	Down	0.000358	0.00056681	AAEL003772-PA OS=Aedes aegypti GN=AAEL003772 PE=4 SV=1
CCG008948.1	981	25	8	1.0950493	0.285082787	-1.941542986	Down	0.000358	0.00056688	AAEL014579-PA OS=Aedes aegypti GN=AAEL014579 PE=3 SV=1
CCG026291.1	524	26	8	2.13208607	0.533714149	-1.998126514	Down	0.0002061	0.0003311	-

CCG025872.1	195	26	8	5.72929795	1.434185714	-1.998126514	Down	0.0002061	0.00033112	AAEL015672-PA OS=Aedes aegypti GN=AAEL015672 PE=4 SV=1
CCG006197.1	391	28	8	3.07711655	0.71525886	-2.105041718	Down	6.68E-05	0.00011095	-
CCG014459.1	288	30	8	4.47601402	0.971063244	-2.204577392	Down	2.11E-05	3.63E-05	AAEL001708-PA OS=Aedes aegypti GN=AAEL001708 PE=4 SV=1
CCG019221.1	624	33	8	2.27243789	0.448183036	-2.342080916	Down	3.58E-06	6.42E-06	Rifesiin heavy chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ015547 PE=3 SV=1
CCG019734.1	558	35	8	2.69523425	0.501193932	-2.426969813	Down	1.07E-06	1.97E-06	AAEL010660-PA OS=Aedes aegypti GN=AAEL010660 PE=3 SV=1
CCG015488.1	558	35	8	2.69523425	0.501193932	-2.426969813	Down	1.07E-06	1.97E-06	AAEL005777-PA OS=Aedes aegypti GN=AAEL005777 PE=4 SV=1
CCG005700.1	1752	36	8	0.88293975	0.159626835	-2.467611798	Down	5.81E-07	1.08E-06	AAEL000751-PA OS=Aedes aegypti GN=AAEL000751 PE=4 SV=1
CCG028245.1	414	37	8	3.84029029	0.675522256	-2.507140162	Down	3.14E-07	5.89E-07	AAEL013513-PA OS=Aedes aegypti GN=AAEL013513 PE=4 SV=1
CCG026699.1	1248	37	8	1.27394245	0.224091518	-2.507140162	Down	3.14E-07	5.89E-07	AAEL008193-PA OS=Aedes aegypti GN=AAEL008193 PE=4 SV=1
CCG019925.1	855	40	8	2.01027998	0.327094987	-2.619614891	Down	4.83E-08	9.40E-08	AAEL010894-PA OS=Aedes aegypti GN=AAEL010894 PE=4 SV=1
CCG015537.1	2399	42	8	0.75228381	0.116576163	-2.690004219	Down	1.36E-08	2.71E-08	AAEL011203-PA OS=Aedes aegypti GN=AAEL011203 PE=4 SV=1
CCG024216.1	963	43	8	1.91869012	0.290411437	-2.723951551	Down	7.20E-09	1.44E-08	-
CCG015837.1	765	43	8	2.41529227	0.365576751	-2.723951551	Down	7.20E-09	1.44E-08	AAEL012315-PA (Fragment) OS=Aedes aegypti GN=AAEL012315 PE=4 SV=1
CCG003489.1	756	46	8	2.61456057	0.369928855	-2.821248752	Down	1.04E-09	2.15E-09	-
CCG001353.1	450	47	8	4.48795006	0.621480476	-2.852275648	Down	5.43E-10	1.13E-09	AAEL015566-PA OS=Aedes aegypti GN=AAEL015566 PE=4 SV=1
CCG012354.1	899	49	8	2.34206562	0.311085889	-2.91239664	Down	1.47E-10	3.12E-10	-
CCG001163.1	5166	53	8	0.44084319	0.05413593	-3.025607251	Down	1.04E-11	2.31E-11	AAEL002227-PA OS=Aedes aegypti GN=AAEL002227 PE=3 SV=1
CCG007152.1	1059	56	8	2.27224281	0.264085188	-3.105041718	Down	1.39E-12	3.36E-12	Protein Wnt (Fragment) OS=Aedes aegypti GN=AAEL014566 PE=3 SV=1

CCG007085.1	948	61	8	2.76493018	0.295006555	-3.228424134	Down	4.67E-14	1.58E-13	AAEL014190-PA OS=Aedes aegypti GN=AAEL014190 PE=4 SV=1
CCG017100.1	1176	62	8	2.26541118	0.237811407	-3.251883107	Down	2.36E-14	8.23E-14	AAEL006980-PA OS=Aedes aegypti GN=AAEL006980 PE=3 SV=1
CCG012817.1	681	64	8	4.03827168	0.410669918	-3.297686796	Down	5.97E-15	2.15E-14	AAEL005540-PA OS=Aedes aegypti GN=AAEL005540 PE=4 SV=1
CCG023590.1	948	73	8	3.30885087	0.295006555	-3.487511355	Down	1.15E-17	4.39E-17	AAEL014190-PA OS=Aedes aegypti GN=AAEL014190 PE=4 SV=1
CCG018257.1	1893	80	8	1.8159423	0.147737039	-3.619614891	Down	8.27E-20	3.22E-19	AAEL009193-PA OS=Aedes aegypti GN=AAEL009193 PE=3 SV=1
CCG002184.1	981	82	8	3.59176171	0.285082787	-3.655238801	Down	2.00E-20	7.83E-20	AAEL007674-PA OS=Aedes aegypti GN=AAEL007674 PE=4 SV=1
CCG012077.1	1311	93	8	3.04819628	0.213322818	-3.836845607	Down	7.57E-24	3.05E-23	AAEL005146-PA OS=Aedes aegypti GN=AAEL005148 PE=4 SV=1
CCG011873.1	318	114	8	15.4042445	0.879453504	-4.13057681	Down	1.71E-30	7.24E-30	-
CCG024590.1	630	128	8	8.73035878	0.443914626	-4.297686796	Down	5.52E-35	2.39E-34	AAEL000359-PA OS=Aedes aegypti GN=AAEL000359 PE=4 SV=1
CCG023498.1	1959	133	8	2.91729183	0.142759681	-4.352969232	Down	1.34E-36	5.88E-36	EV-riano domain-containing protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ015374 PE=4 SV=1
CCG017792.1	912	171	8	8.05682524	0.306651551	-4.715539311	Down	5.46E-49	2.53E-48	AAEL005989-PA OS=Aedes aegypti GN=AAEL005989 PE=4 SV=1
CCG017947.1	1098	196	8	7.67037157	0.254705113	-4.91239664	Down	3.10E-57	1.48E-56	Beta-1,5-N-acetylglucosaminyl transferase radical fringe OS=Culex quinquefasciatus GN=CpipJ_CPIJ010821 PE=4 SV=1
CCG026543.1	1002	250	8	10.7209917	0.279107998	-5.263471081	Down	3.26E-75	1.64E-74	AAEL002716-PA OS=Aedes aegypti GN=AAEL002716 PE=4 SV=1
CCG021979.1	1119	362	8	13.9008435	0.249925124	-5.797532683	Down	6.04E-113	3.26E-112	AAEL011591-PA OS=Aedes aegypti GN=AAEL011591 PE=4 SV=1
CCG004605.1	729	372	8	21.9269428	0.383629923	-5.836845607	Down	2.47E-116	1.34E-115	AAEL008697-PA OS=Aedes aegypti GN=AAEL008697 PE=4 SV=1
CCG014841.1	1314	382	8	12.4919624	0.212835779	-5.875115624	Down	1.00E-119	5.50E-119	AGAP005519-PA OS=Anopheles gambiae GN=AGAP005519 PE=4 SV=3

CCG020506.1	1908	514	8	11.5757042	0.146575584	-6.303311345	Down	1.16E-164	6.69E-164	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018057 PE=4 SV=1
CCG004325.1	1278	673	8	22.6280371	0.218831154	-6.692149491	Down	4.32E-219	2.61E-218	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000931 PE=4 SV=1
CCG017334.1	690	727	8	45.2739088	0.405313354	-6.80349835	Down	1.26E-237	7.72E-237	40S ribosomal protein S13 OS=Aedes aegypti GN=40sRpS13 PE=2 SV=1
CCG025653.1	231	1283	8	238.658742	1.210676252	-7.622992251	Down	0	0	AAEL006011-FA OS=Aedes aegypti GN=AAEL006011 PE=4 SV=1
CCG016568.1	519	1800	8	149.027981	0.538855904	-8.111467987	Down	0	0	AAEL010101-FA OS=Aedes aegypti GN=AAEL010161 PE=4 SV=1
CCG017457.1	936	2958	8	135.795379	0.29878869	-8.828093133	Down	0	0	AAEL010104-FA OS=Aedes aegypti GN=AAEL010164 PE=4 SV=1
CCG026411.1	1878	5499	8	125.820325	0.148917047	-9.722640367	Down	0	0	F-box/inr protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011368 PE=4 SV=1
CCG014543.1	1005	26	9	1.11165483	0.313059195	-1.828201513	Down	0.0004711	0.00074144	AAEL015307-FA OS=Aedes aegypti GN=AAEL015307 PE=4 SV=1
CCG004217.1	2094	26	9	0.53353061	0.150250473	-1.828201513	Down	0.0004711	0.00074148	AAEL007588-FA OS=Aedes aegypti GN=AAEL007588 PE=4 SV=1
CCG019642.1	1431	27	9	0.81074971	0.219863376	-1.882649297	Down	0.0002763	0.00044048	Retrovirus-related Pol polyprotein from transposon TNT 1-94 OS=Trichinella spiralis GN=Tsp_01017 PE=4 SV=1
CCG001088.1	386	27	9	3.00565501	0.815089355	-1.882649297	Down	0.0002763	0.00044051	-
CCG011797.1	1056	28	9	1.13934902	0.297939859	-1.935116717	Down	0.0001608	0.00025947	-
CCG012526.1	608	28	9	1.97886936	0.517474492	-1.935116717	Down	0.0001608	0.00025948	-
CCG018451.1	545	29	9	2.28646294	0.577292644	-1.98574279	Down	9.29E-05	0.00015302	-
CCG026510.1	858	29	9	1.452357	0.366695211	-1.98574279	Down	9.29E-05	0.00015303	AAEL010894-FA OS=Aedes aegypti GN=AAEL010894 PE=4 SV=1
CCG001820.1	657	31	9	2.02749128	0.478880504	-2.081958105	Down	3.04E-05	5.16E-05	-
CCG025487.1	1008	32	9	1.36411856	0.312127471	-2.127761795	Down	1.72E-05	2.97E-05	AAEL000806-FA OS=Aedes aegypti GN=AAEL000806 PE=4 SV=1

CCG013184.1	783	33	9	1.81098498	0.401819273	-2.172155914	Down	9.70E-06	1.70E-05	AAEL003642-PA OS=Reedes aegypti GN=AAEL003642 PE=3 SV=1
CCG005977.1	942	33	9	1.50530917	0.333996275	-2.172155914	Down	9.70E-06	1.70E-05	AAEL001318-PA OS=Reedes aegypti GN=AAEL001318 PE=4 SV=1
CCG018828.1	840	34	9	1.73925116	0.374552965	-2.215224636	Down	5.43E-06	9.65E-06	AAEL011530-PA OS=Reedes aegypti GN=AAEL011530 PE=4 SV=1
CCG001394.1	1437	35	9	1.04658365	0.218945366	-2.257044812	Down	3.03E-06	5.44E-06	AAEL008715-PA OS=Reedes aegypti GN=AAEL008715 PE=4 SV=1
CCG006388.1	522	35	9	2.88111247	0.60272891	-2.257044812	Down	3.03E-06	5.44E-06	AAEL005264-PA OS=Reedes aegypti GN=AAEL005264 PE=4 SV=1
CCG006481.1	291	36	9	5.31584346	1.081183818	-2.297686796	Down	1.68E-06	3.06E-06	Vesicular monoamine transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ007766 PE=4 SV=1
CCG000029.1	522	36	9	2.96342997	0.60272891	-2.297686796	Down	1.68E-06	3.06E-06	AAEL010040-PA OS=Reedes aegypti GN=AAEL010040 PE=4 SV=1
CCG002642.1	1496	36	9	1.03403105	0.210310489	-2.297686796	Down	1.68E-06	3.06E-06	AAEL014435-PA OS=Reedes aegypti GN=AAEL014435 PE=4 SV=1
CCG009590.1	597	37	9	2.66311588	0.527009198	-2.33721516	Down	9.26E-07	1.71E-06	-
CCG013796.1	1139	38	9	1.43358202	0.276228701	-2.375689308	Down	5.09E-07	9.49E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010109 PE=4 SV=1
CCG025372.1	603	40	9	2.85039699	0.521765325	-2.44968989	Down	1.51E-07	2.88E-07	AAEL012052-PA OS=Reedes aegypti GN=AAEL012052 PE=4 SV=1
CCG016348.1	864	43	9	2.13854003	0.364148716	-2.554026549	Down	2.39E-08	4.69E-08	AAEL008471-PA OS=Reedes aegypti GN=AAEL008471 PE=4 SV=1
CCG001958.1	505	50	9	4.25442917	0.623018794	-2.771617985	Down	2.84E-10	5.97E-10	-
CCG006432.1	534	51	9	4.10385106	0.58918444	-2.800187137	Down	1.49E-10	3.17E-10	AAEL009333-PA OS=Reedes aegypti GN=AAEL009333 PE=4 SV=1
CCG025242.1	486	55	9	4.86283005	0.647375496	-2.909121508	Down	1.10E-11	2.44E-11	AAEL009744-PA OS=Reedes aegypti GN=AAEL009744 PE=4 SV=1
CCG009089.1	396	59	9	6.40205642	0.79450629	-3.010404844	Down	7.83E-13	1.96E-12	AAEL010061-PA OS=Reedes aegypti GN=AAEL010061 PE=4 SV=1
CCG019645.1	656	63	9	4.12666659	0.479610504	-3.105041718	Down	5.39E-14	1.80E-13	Uncharacterized protein OS=Clonorchis sinensis GN=CLF_106999 PE=4 SV=1

CCG024708.1	1614	72	9	1.91686548	0.194934629	-3.297686796	Down	1.18E-16	4.44E-16	AAEL002670-PA OS=Aedes aegypti GN=AAEL002670 PE=4 SV=1
CCG002587.1	3975	83	9	0.89722968	0.079150815	-3.502801226	Down	5.65E-20	2.20E-19	AAEL003262-PA OS=Aedes aegypti GN=AAEL003262 PE=4 SV=1
CCG009617.1	1116	89	9	3.42679783	0.281921587	-3.603495226	Down	8.24E-22	3.26E-21	AAEL009631-PA OS=Aedes aegypti GN=AAEL009631 PE=3 SV=1
CCG028595.1	486	91	9	8.04577335	0.647375496	-3.635556435	Down	2.00E-22	7.95E-22	AAEL008450-PA OS=Aedes aegypti GN=AAEL008450 PE=4 SV=1
CCG010275.1	987	97	9	4.22296277	0.318768481	-3.727674637	Down	2.78E-24	1.12E-23	AAEL011633-PA OS=Aedes aegypti GN=AAEL011633 PE=4 SV=1
CCG016054.1	1111	98	9	3.79030962	0.283190361	-3.742471639	Down	1.36E-24	5.50E-24	Yokozuna protein OS=Bombyx mori GN=Yokozuna PE=4 SV=1
CCG003860.1	619	119	9	8.26074058	0.508278661	-4.022579558	Down	3.43E-31	1.46E-30	-
CCG020463.1	1020	127	9	5.35015323	0.308455383	-4.116446482	Down	9.80E-34	4.21E-33	AAEL011459-PA OS=Aedes aegypti GN=AAEL011459 PE=4 SV=1
CCG028621.1	708	139	9	8.43614846	0.444384874	-4.246702867	Down	1.41E-37	6.21E-37	AAEL014235-PA OS=Aedes aegypti GN=AAEL014235 PE=4 SV=1
CCG012104.1	1164	140	9	5.16818114	0.270295954	-4.257044812	Down	6.72E-38	2.97E-37	Serine protease inhibitor OS=Culex quinquefasciatus GN=CpipJ_CPIJ006815 PE=3 SV=1
CCG009158.1	1209	142	9	5.04690018	0.260235311	-4.277508914	Down	1.53E-38	6.78E-38	Lipase OS=Aedes aegypti GN=AAEL012349 PE=3 SV=1
CCG001894.1	1230	145	9	5.06553782	0.255792269	-4.307670885	Down	1.65E-39	7.35E-39	AAEL010369-PA OS=Aedes aegypti GN=AAEL010369 PE=4 SV=1
CCG009533.1	462	149	9	13.8582045	0.681005392	-4.346930315	Down	8.45E-41	3.78E-40	-
CCG006633.1	867	150	9	7.43421014	0.362888686	-4.356580485	Down	4.02E-41	1.80E-40	AgCP7521-like protein (Fragment) OS=Aedes albopictus PE=4 SV=1
CCG006994.1	517	157	9	13.0488362	0.60855801	-4.422382544	Down	2.17E-43	9.84E-43	AAEL007416-PA OS=Aedes aegypti GN=AAEL007416 PE=4 SV=1
CCG027454.1	705	177	9	10.7881461	0.446275874	-4.595367345	Down	6.68E-50	3.11E-49	AAEL003216-PA OS=Aedes aegypti GN=AAEL003216 PE=4 SV=1
CCG025426.1	837	178	9	9.13812755	0.375895449	-4.603495226	Down	3.15E-50	1.47E-49	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007357 PE=4 SV=1

CCG011146.1	1488	187	9	5.40009434	0.21144119	-4.674656255	Down	3.55E-53	1.68E-52	AAEL011310-PA OS=Aedes aegypti GN=AAEL011310 PE=4 SV=1
CCG021307.1	573	243	9	18.222767	0.549082881	-5.052574298	Down	1.11E-71	5.53E-71	AAEL001290-PA OS=Aedes aegypti GN=AAEL001290 PE=4 SV=1
CCG013754.1	807	249	9	13.2583196	0.389869258	-5.087763727	Down	1.12E-73	5.62E-73	AAEL011200-PA OS=Aedes aegypti GN=AAEL011206 PE=4 SV=1
CCG025465.1	422	464	9	47.2463433	0.745555666	-5.98574279	Down	3.83E-146	2.17E-145	AAEL007410-PA OS=Aedes aegypti GN=AAEL007416 PE=4 SV=1
CCG016952.1	918	652	9	30.5188093	0.342728204	-6.476489949	Down	2.79E-210	1.67E-209	AAEL007434-PA OS=Aedes aegypti GN=AAEL007434 PE=4 SV=1
CCG024532.1	993	1025	9	44.3544592	0.316842388	-7.129169989	Down	0	0	AAEL012770-PA OS=Aedes aegypti GN=CYP325N1 PE=3 SV=2
CCG016528.1	2649	1235	9	20.0330775	0.118771042	-7.398057121	Down	0	0	AAEL008496-PA OS=Aedes aegypti GN=AAEL008498 PE=4 SV=1
CCG017456.1	477	1631	9	146.925864	0.659590128	-7.799302861	Down	0	0	AAEL010180-PA OS=Aedes aegypti GN=AAEL010186 PE=4 SV=1
CCG027516.1	456	4308	9	405.950914	0.689965989	-9.200564329	Down	0	0	-
CCG014501.1	384	27	10	3.02130946	0.910371791	-1.730646203	Down	0.0005996	0.00093314	AAEL006355-PA OS=Aedes aegypti GN=SCRC1 PE=4 SV=2
CCG002905.1	1056	28	10	1.13934902	0.331044288	-1.783113623	Down	0.0003576	0.0005665	AAEL009966-PA OS=Aedes aegypti GN=AAEL009966 PE=4 SV=1
CCG012198.1	1359	28	10	0.88532198	0.257235296	-1.783113623	Down	0.0003576	0.00056654	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008569 PE=4 SV=1
CCG021640.1	1662	28	10	0.72391851	0.210338609	-1.783113623	Down	0.0003576	0.00056647	AGAP007672-PA OS=Anopheles gambiae GN=AgaP_AGAP007672 PE=3 SV=5
CCG016745.1	1026	28	10	1.17266332	0.340723945	-1.783113623	Down	0.0003576	0.0005664	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009588 PE=4 SV=1
CCG005844.1	1202	28	10	1.00095888	0.290834249	-1.783113623	Down	0.0003576	0.00056643	Uncharacterized protein OS=Anopheles darlingi GN=AND_05226 PE=4 SV=1

CCG021260.1	1852	28	10	0.64965042	0.188759594	-1.783113623	Down	0.0003576	0.00056636	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC006955 PE=4 SV=1
CCG002182.1	1497	29	10	0.83241303	0.233522223	-1.833739696	Down	0.0002117	0.00033973	AAEL007675-PA (Fragment) OS=Aedes aegypti GN=AAEL007675 PE=3 SV=1
CCG004073.1	750	29	10	1.6614964	0.466110357	-1.833739696	Down	0.0002117	0.00033975	AAEL002919-PA OS=Aedes aegypti GN=AAEL002919 PE=4 SV=1
CCG004742.1	210	36	10	7.36624022	1.664679846	-2.145683703	Down	4.46E-06	7.95E-06	AAEL000711-PA OS=Aedes aegypti GN=AAEL000711 PE=4 SV=1
CCG009578.1	409	36	10	3.78217713	0.854725593	-2.145683703	Down	4.46E-06	7.95E-06	-
CCG016597.1	747	37	10	2.12835366	0.467982286	-2.185212067	Down	2.51E-06	4.53E-06	Uncharacterized protein OS=Anopheles darlingi GN=AND_17075 PE=4 SV=1
CCG019712.1	534	38	10	3.05777138	0.654649378	-2.223686215	Down	1.41E-06	2.57E-06	AAEL014505-PA OS=Aedes aegypti GN=AAEL014505 PE=2 SV=1
CCG002242.1	333	40	10	5.16152968	1.049798101	-2.297686796	Down	4.36E-07	8.14E-07	-
CCG004031.1	229	46	10	8.63147507	1.526562304	-2.499320657	Down	1.17E-08	2.32E-08	-
CCG025544.1	360	50	10	5.9680187	0.971063244	-2.619614891	Down	9.72E-10	2.01E-09	Voltage-dependent para-like sodium channel (Fragment) OS=Aedes albopictus PE=2 SV=1
CCG020241.1	1088	53	10	2.09319479	0.321307691	-2.703679156	Down	1.46E-10	3.11E-10	AAEL007063-PA OS=Aedes aegypti GN=AAEL007063 PE=3 SV=1
CCG003675.1	1659	63	10	1.63176207	0.210718968	-2.953038625	Down	2.23E-13	6.38E-13	AAEL013194-PA OS=Aedes aegypti GN=AAEL013194 PE=4 SV=1
CCG027003.1	822	69	10	3.60694853	0.425283172	-3.084283158	Down	4.11E-15	1.49E-14	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003408 PE=4 SV=1
CCG013311.1	807	84	10	4.47268613	0.433188064	-3.368076124	Down	1.48E-19	5.75E-19	AAEL013621-PA OS=Aedes aegypti GN=AAEL013621 PE=4 SV=1
CCG009727.1	1926	89	10	1.98562117	0.181507148	-3.451492132	Down	4.58E-21	1.81E-20	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013977 PE=4 SV=1

CCG014067.1	2043	92	10	1.93500518	0.171112466	-3.499320657	Down	5.62E-22	2.23E-21	Pro-resilin OS=Culex quinquefasciatus GN=CpipJ_CPIJ001605 PE=4 SV=1
CCG024763.1	657	104	10	6.80190624	0.532089449	-3.676198419	Down	1.17E-25	4.77E-25	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010720 PE=4 SV=1
CCG019833.1	1110	105	10	4.06470463	0.31493943	-3.690004219	Down	5.73E-26	2.35E-25	AAEL014938-PA OS=Aedes aegypti GN=AAEL014938 PE=3 SV=1
CCG020293.1	363	123	10	14.5599927	0.963037928	-3.918273207	Down	1.37E-31	5.83E-31	AAEL017144-PA OS=Aedes aegypti PE=4 SV=1
CCG002220.1	543	156	10	12.3448961	0.643798836	-4.26116092	Down	4.25E-42	1.91E-41	AAEL010346-PA OS=Aedes aegypti GN=AAEL010346 PE=4 SV=1
CCG021406.1	2694	212	10	3.3814342	0.129763462	-4.703679156	Down	2.58E-60	1.25E-59	AAEL005240-PA OS=Aedes aegypti GN=AAEL005240 PE=4 SV=1
CCG012989.1	1125	226	10	8.63214224	0.310740238	-4.795937664	Down	6.41E-65	3.13E-64	AAEL005621-PA OS=Aedes aegypti GN=GPROP5 PE=3 SV=1
CCG008100.1	1407	289	10	8.82605068	0.248459679	-5.150684384	Down	8.17E-86	4.20E-85	AAEL011375-PA OS=Aedes aegypti GN=AAEL011375 PE=3 SV=1
CCG006228.1	1734	376	10	9.31754338	0.201604826	-5.530347553	Down	5.51E-115	2.99E-114	AAEL011774-PA OS=Aedes aegypti GN=AAEL011774 PE=4 SV=1
CCG017394.1	2847	856	10	12.9195971	0.122789873	-6.717225688	Down	1.44E-278	8.98E-278	AAEL006763-PA OS=Aedes aegypti GN=AAEL006763 PE=4 SV=1
CCG016157.1	591	1275	10	92.7012041	0.591510605	-7.292040233	Down	0	0	AGAP006829-PA OS=Anopheles gambiae GN=CPR59 PE=4 SV=4
CCG016889.1	1920	2686	10	60.1128683	0.182074358	-8.367002291	Down	0	0	AAEL009910-PA OS=Aedes aegypti GN=AAEL009910 PE=4 SV=1
CCG016890.1	900	4632	10	221.150901	0.388425297	-9.153178239	Down	0	0	AAEL009904-PA OS=Aedes aegypti GN=AAEL009904 PE=4 SV=1
CCG006366.1	1131	34833	10	1323.39944	0.309091749	-12.06392581	Down	0	0	AAEL009451-PA OS=Aedes aegypti GN=AAEL009451 PE=3 SV=1
CCG013365.1	1881	29	11	0.66247863	0.204434367	-1.696236173	Down	0.0004498	0.00070891	AAEL013026-PA (Fragment) OS=Aedes aegypti GN=AAEL013026 PE=3 SV=1

CCG002058.1	924	30	11	1.39512125	0.416169962	-1.745145773	Down	0.0002703	0.00043136	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007764 PE=4 SV_1
CCG003719.1	283	34	11	5.16244161	1.358802277	-1.925718019	Down	3.29E-05	5.57E-05	-
CCG019107.1	474	34	11	3.08221725	0.811268026	-1.925718019	Down	3.29E-05	5.57E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008218 PE=4 SV_1
CCG021701.1	1077	35	11	1.39641663	0.357048324	-1.967538194	Down	1.91E-05	3.28E-05	Translation elongation factor EF-1 alpha/Tu OS=Culex quinquefasciatus GN=CpipJ_CPIJ010941 PE=4 SV_1
CCG012190.1	789	36	11	1.96059626	0.48737775	-2.008180179	Down	1.10E-05	1.92E-05	AAEL009233-PA (Fragment) OS=Aedes aegypti GN=AAEL009233 PE=4 SV=1
CCG001469.1	1591	36	11	0.97228815	0.241697702	-2.008180179	Down	1.10E-05	1.93E-05	AAEL009966-PA OS=Aedes aegypti GN=AAEL009966 PE=4 SV_1
CCG023394.1	1539	36	11	1.00513999	0.249864226	-2.008180179	Down	1.10E-05	1.93E-05	AAEL006846-PA OS=Aedes aegypti GN=AAEL006846 PE=4 SV_1
CCG011348.1	1212	41	11	1.45359663	0.31727809	-2.195807182	Down	6.54E-07	1.21E-06	AAEL004174-PA OS=Aedes aegypti GN=AAEL004174 PE=4 SV_1
CCG015828.1	1458	42	11	1.23781129	0.263745572	-2.2305726	Down	3.66E-07	6.87E-07	AAEL008336-PA (Fragment) OS=Aedes aegypti GN=AAEL008336 PE=4 SV=1
CCG018005.1	777	42	11	2.32268836	0.494904819	-2.2305726	Down	3.66E-07	6.87E-07	AAEL004525-PA OS=Aedes aegypti GN=AAEL004525 PE=4 SV_1
CCG014098.1	498	43	11	3.71023813	0.772170772	-2.264519932	Down	2.04E-07	3.87E-07	AAEL004627-PA OS=Aedes aegypti GN=AAEL004627 PE=4 SV_1
CCG027837.1	363	45	11	5.3268266	1.05934172	-2.330108274	Down	6.28E-08	1.22E-07	AAEL017144-PA OS=Aedes aegypti PE=4 SV=1
CCG010055.1	339	45	11	5.70394707	1.134339364	-2.330108274	Down	6.28E-08	1.22E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008247 PE=4 SV_1
CCG001391.1	807	46	11	2.44932812	0.476506871	-2.361817134	Down	3.46E-08	6.76E-08	cGMP-dependent protein kinase OS=Aedes aegypti GN=AAEL007826 PE=3 SV=1

CCG024997.1	456	51	11	4.80582558	0.843291764	-2.51068052	Down	1.67E-09	3.41E-09	AAEL002626-PA OS=Aedes aegypti GN=AAEL002626 PE=4 SV=1
CCG024241.1	627	51	11	3.49514588	0.613303101	-2.51068052	Down	1.67E-09	3.41E-09	AAEL002839-PA (Fragment) OS=Aedes aegypti GN=AAEL002839 PE=4 SV=1
CCG027162.1	768	51	11	2.85345894	0.500704485	-2.51068052	Down	1.67E-09	3.41E-09	AAEL012065-PA OS=Aedes aegypti GN=AAEL012065 PE=4 SV=1
CCG006669.1	3183	54	11	0.72898702	0.120810884	-2.59314268	Down	2.60E-10	5.47E-10	AAEL010252-PA OS=Aedes aegypti GN=AAEL010252 PE=4 SV=1
CCG000208.1	1915	54	11	1.2116792	0.200804723	-2.59314268	Down	2.60E-10	5.47E-10	Putative finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014131 PE=4 SV=1
CCG023554.1	1320	60	11	1.95316976	0.291318973	-2.745145773	Down	5.85E-12	1.32E-11	-
CCG017022.1	578	62	11	4.60921029	0.665295925	-2.792451488	Down	1.62E-12	3.89E-12	-
CCG000580.1	1401	63	11	1.93225787	0.27447612	-2.815535101	Down	8.50E-13	2.12E-12	Cadherin EGF LAG seven-pass G-type receptor 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002422 PE=4 SV=1
CCG022514.1	666	64	11	4.12922375	0.577388956	-2.838255178	Down	4.45E-13	1.17E-12	AAEL014418-PA OS=Aedes aegypti GN=AAEL014418 PE=4 SV=1
CCG018054.1	303	65	11	9.21792987	1.269112358	-2.860622991	Down	2.32E-13	6.62E-13	AAEL011332-PA OS=Aedes aegypti GN=AAEL011332 PE=4 SV=1
CCG026535.1	963	68	11	3.03420764	0.399315726	-2.925718019	Down	3.27E-14	1.13E-13	AAEL002716-PA OS=Aedes aegypti GN=AAEL002716 PE=4 SV=1
CCG009324.1	1632	69	11	1.8167351	0.23562564	-2.946779634	Down	1.69E-14	5.97E-14	AAEL008129-PA OS=Aedes aegypti GN=AAEL008129 PE=4 SV=1
CCG011777.1	219	88	11	17.2663774	1.75589518	-3.297686796	Down	4.66E-20	1.82E-19	AAEL011768-PA OS=Aedes aegypti GN=AAEL011768 PE=4 SV=1
CCG022049.1	1417	97	11	2.94147089	0.271376884	-3.43816802	Down	9.20E-23	3.67E-22	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004282 PE=4 SV=1
CCG012810.1	3279	103	11	1.34976599	0.117273878	-3.524755705	Down	1.38E-24	5.60E-24	Uncharacterized protein OS=Anopheles darlingi GN=AND_23101 PE=4 SV=1

CCG026967.1	2130	106	11	2.13839994	0.180535702	-3.566175632	Down	1.67E-25	6.84E-25	Imp4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017542 PE=4 SV=1
CCG005261.1	489	114	11	10.0174841	0.786382504	-3.671145192	Down	5.80E-28	2.41E-27	AAEL008765-PA OS=Aedes aegypti GN=AAEL008765 PE=4 SV=1
CCG024294.1	1611	127	11	3.38743408	0.23869711	-3.826939864	Down	5.27E-32	2.24E-31	AAEL007582-PA OS=Aedes aegypti GN=AAEL007582 PE=3 SV=1
CCG005575.1	300	164	11	23.4901216	1.281803482	-4.195807182	Down	9.96E-44	4.52E-43	Uncharacterized protein OS=Anopheles darlingi GN=AND_27279 PE=4 SV=1
CCG010610.1	1626	169	11	4.46610403	0.236495107	-4.239134614	Down	2.48E-45	1.13E-44	AAEL003426-PA (Fragment) OS=Aedes aegypti GN=AAEL003426 PE=4 SV=1
CCG012476.1	943	178	11	8.11093612	0.407784777	-4.313988609	Down	3.15E-48	1.46E-47	AAEL015464-PA OS=Aedes aegypti GN=AAEL015464 PE=3 SV=1
CCG007339.1	1833	201	11	4.71190216	0.209787804	-4.489306869	Down	1.12E-55	5.32E-55	Putative Ropn11-like protein OS=Anopheles gambiae PE=2 SV=1
CCG001442.1	918	211	11	9.87648584	0.418890027	-4.559354366	Down	6.18E-59	2.96E-58	AAEL003216-PA OS=Aedes aegypti GN=AAEL003216 PE=4 SV=1
CCG012988.1	1125	404	11	15.4309091	0.341814262	-5.49646666	Down	4.17E-123	2.29E-122	AAEL005625-PA OS=Aedes aegypti GN=GPROP4 PE=3 SV=1
CCG006836.1	1062	505	11	20.4328776	0.362091379	-5.818394755	Down	3.20E-157	1.83E-156	AAEL001745-PA OS=Aedes aegypti GN=AAEL001745 PE=4 SV=1
CCG019360.1	432	505	11	50.230824	0.890141307	-5.818394755	Down	3.20E-157	1.83E-156	AAEL008681-PA OS=Aedes aegypti GN=CTL12 PE=4 SV=1
CCG014523.1	621	2769	11	191.599348	0.619228735	-8.273404516	Down	0	0	AAEL010824-PA OS=Aedes aegypti GN=AAEL010824 PE=4 SV=1
CCG005687.1	1407	30	12	0.91619903	0.298151614	-1.619614891	Down	0.0005521	0.0008649	AAEL009592-PA OS=Aedes aegypti GN=AAEL009592 PE=4 SV=1
CCG000656.1	329	32	12	4.17942707	1.275073925	-1.712724295	Down	0.0002037	0.00032732	Cecropin A OS=Culex quinquefasciatus GN=CpipJ_CPIJ010699 PE=4 SV=1
CCG006655.1	456	33	12	3.10965185	0.919954652	-1.757118415	Down	0.0001224	0.0001989	AAEL008573-PA (Fragment) OS=Aedes aegypti GN=AAEL008573 PE=4 SV=1

CCG011871.1	1339	33	12	1.05900018	0.313292996	-1.757118415	Down	0.0001224	0.00019891	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001097 PE=4 SV=1
CCG009455.1	674	38	12	2.42262599	0.622402554	-1.960651809	Down	8.69E-06	1.52E-05	-
CCG028205.1	1701	38	12	0.95993528	0.246619236	-1.960651809	Down	8.69E-06	1.52E-05	AAEL013023-PA OS=Aedes aegypti GN=AAEL013023 PE=4 SV=1
CCG007225.1	618	40	12	2.7812126	0.678801491	-2.03465239	Down	2.90E-06	5.21E-06	AAEL012571-PA OS=Aedes aegypti GN=AAEL012571 PE=4 SV=1
CCG028402.1	849	41	12	2.07509908	0.494109919	-2.0702763	Down	1.66E-06	3.03E-06	AgCP7521-like protein (Fragment) OS=Aedes albopictus PE=4 SV=1
CCG014369.1	687	41	12	2.56442375	0.610624922	-2.0702763	Down	1.66E-06	3.03E-06	AAEL014864-PA OS=Aedes aegypti GN=AAEL014864 PE=4 SV=1
CCG015660.1	1087	42	12	1.66028413	0.385923939	-2.105041718	Down	9.47E-07	1.75E-06	-
CCG023872.1	573	43	12	3.22460487	0.732110508	-2.13898905	Down	5.37E-07	1.00E-06	AAEL011492-PA OS=Aedes aegypti GN=AAEL011492 PE=4 SV=1
CCG014181.1	1593	47	12	1.2677825	0.263339185	-2.267313147	Down	5.34E-08	1.04E-07	AAEL002623-PA OS=Aedes aegypti GN=AAEL002623 PE=4 SV=1
CCG016075.1	957	54	12	2.42462452	0.438348298	-2.467611798	Down	8.13E-10	1.68E-09	Zinc carboxypeptidase OS=Culex quinquefasciatus GN=CpipJ_CPIJ010796 PE=4 SV=1
CCG005895.1	762	57	12	3.21427149	0.550524044	-2.54561431	Down	1.29E-10	2.75E-10	AAEL002954-PA OS=Aedes aegypti GN=AAEL002954 PE=4 SV=1
CCG003222.1	1893	60	12	1.36195672	0.221605558	-2.619614891	Down	1.99E-11	4.36E-11	AAEL008325-PA OS=Aedes aegypti GN=AAEL008325 PE=4 SV=1
CCG026997.1	378	73	12	8.2983879	1.109786564	-2.902548854	Down	4.79E-15	1.73E-14	AAEL017144-PA OS=Aedes aegypti PE=4 SV=1
CCG017612.1	726	75	12	4.43902217	0.577822757	-2.941542986	Down	1.29E-15	4.69E-15	Probable transport protein Sec61 alpha subunit OS=Aedes aegypti GN=Sec61 PE=2 SV=1
CCG022339.1	396	80	12	8.68075447	1.05934172	-3.03465239	Down	4.71E-17	1.78E-16	-
CCG003722.1	1416	83	12	2.5187062	0.296256583	-3.087763727	Down	6.35E-18	2.42E-17	AAEL007806-PA OS=Aedes aegypti GN=CYP4D39 PE=3 SV=1
CCG022575.1	442	89	12	8.65227688	0.949093487	-3.188457726	Down	1.11E-19	4.31E-19	-
CCG025138.1	534	105	12	8.44910512	0.785579253	-3.426969813	Down	1.85E-24	7.49E-24	AAEL011642-PA OS=Aedes aegypti GN=AAEL011642 PE=4 SV=1

CCG007231.1	1743	114	12	2.81041293	0.240676604	-3.54561431	Down	3.41E-27	1.41E-26	AAEL010201-PA OS=Aedes aegypti GN=AAEL010201 PE=4 SV=1
CCG026150.1	2646	115	12	1.86754327	0.158540938	-3.558214346	Down	1.69E-27	6.98E-27	AAEL000778-PA OS=Aedes aegypti GN=AAEL000778 PE=4 SV=1
CCG023260.1	1275	142	12	4.78564887	0.329019076	-3.862471415	Down	7.22E-36	3.15E-35	AAEL0006994-PA OS=Aedes aegypti GN=AAEL006994 PE=4 SV=1
CCG007287.1	2055	145	12	3.03192775	0.204135923	-3.892633385	Down	8.26E-37	3.62E-36	AAEL000247-PA OS=Aedes aegypti GN=AAEL000247 PE=4 SV=1
CCG016642.1	2024	155	12	3.29066644	0.207262511	-3.988848701	Down	5.82E-40	2.60E-39	AAEL0009966-PA OS=Aedes aegypti GN=AAEL009966 PE=4 SV=1
CCG013211.1	648	157	12	10.4108771	0.647375496	-4.007345044	Down	1.35E-40	6.06E-40	AAEL005770-PA OS=Aedes aegypti GN=AAEL005770 PE=4 SV=1
CCG000082.1	1779	166	12	4.00954241	0.235806251	-4.087763727	Down	1.88E-43	8.53E-43	AAEL001835-PA OS=Aedes aegypti GN=AAEL001835 PE=3 SV=1
CCG025094.1	1335	246	12	7.91801851	0.314231701	-4.655238801	Down	2.46E-69	1.22E-68	Ams protein transcript variant 1 (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG022279.1	3624	580	12	6.87705466	0.115755883	-5.892633385	Down	3.04E-181	1.78E-180	AAEL005967-PA OS=Aedes aegypti GN=AAEL005967 PE=4 SV=1
CCG018643.1	711	748	12	45.205853	0.59001311	-6.259618755	Down	2.01E-238	1.23E-237	AAEL009577-PA OS=Aedes aegypti GN=AAEL009577 PE=4 SV=1
CCG017802.1	351	931	12	113.973854	1.195154761	-6.575361653	Down	5.28E-301	3.33E-300	-
CCG011672.1	483	2087	12	185.668398	0.868528615	-7.739939181	Down	0	0	-
CCG025882.1	348	0	13	0.001	1.305912638	10.35084267	Up	0.0004804	0.00075505	AAEL013443-PA OS=Aedes aegypti GN=AAEL013443 PE=4 SV=1
CCG009113.1	445	0	13	0.001	1.021253029	9.996124642	Up	0.0004804	0.00075556	-
CCG001329.1	462	0	13	0.001	0.983674455	9.942037127	Up	0.0004804	0.00075585	AAEL004611-PA OS=Aedes aegypti GN=AAEL004611 PE=4 SV=1
CCG017910.1	465	0	13	0.001	0.977328168	9.932699262	Up	0.0004804	0.00075523	AAEL004145-PA OS=Aedes aegypti GN=AAEL004145 PE=4 SV=1
CCG010936.1	486	0	13	0.001	0.935097938	9.868973665	Up	0.0004804	0.0007558	-
CCG008688.1	507	0	13	0.001	0.896366071	9.807944231	Up	0.0004804	0.00075566	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 SV=1
CCG002798.1	538	0	13	0.001	0.844716725	9.722323806	Up	0.0004804	0.00075528	-

CCG009511.1	600	0	13	0.001	0.75742933	9.564967478	Up	0.0004804	0.000755	AAEL001383-PA OS=Aedes aegypti GN=AAEL001383 PE=4 SV=1
CCG022386.1	604	0	13	0.001	0.752413242	9.555381429	Up	0.0004804	0.00075547	-
CCG019765.1	693	0	13	0.001	0.65578297	9.357074626	Up	0.0004804	0.00075538	AAEL004751-PA OS=Aedes aegypti GN=AAEL004751 PE=4 SV=1
CCG009150.1	744	0	13	0.001	0.610830105	9.254627357	Up	0.0004804	0.00075514	-
CCG025268.1	972	0	13	0.001	0.467548969	8.868973665	Up	0.0004804	0.00075561	AAEL007999-PA OS=Aedes aegypti GN=AAEL007999 PE=4 SV=1
CCG003137.1	1032	0	13	0.001	0.44036589	8.782558913	Up	0.0004804	0.00075571	AAEL015565-PA OS=Aedes aegypti GN=AAEL015565 PE=4 SV=1
CCG021419.1	1152	0	13	0.001	0.394494443	8.623861167	Up	0.0004804	0.00075575	AAEL010973-PA OS=Aedes aegypti GN=AAEL010973 PE=4 SV=1
CCG019144.1	1170	0	13	0.001	0.388425297	8.601493354	Up	0.0004804	0.00075589	AAEL006371-PA OS=Aedes aegypti GN=AAEL006371 PE=3 SV=1
CCG025364.1	1593	0	13	0.001	0.285284117	8.156255617	Up	0.0004804	0.00075519	AAEL009243-PA OS=Aedes aegypti GN=AAEL009243 PE=3 SV=1
CCG011001.1	1617	0	13	0.001	0.281049844	8.134682205	Up	0.0004804	0.00075509	Putative uncharacterized protein GLEAN_15470 OS=Tribolium castaneum GN=GLEAN_15470 PE=4 SV=1
CCG004788.1	1896	0	13	0.001	0.239692826	7.905042919	Up	0.0004804	0.00075552	-
CCG006667.1	2652	0	13	0.001	0.171364102	7.420921108	Up	0.0004804	0.00075542	Uncharacterized protein OS=Anopheles darlingi GN=AND_19618 PE=4 SV=1
CCG007445.1	2793	0	13	0.001	0.162713068	7.34618631	Up	0.0004804	0.00075533	AAEL006287-PA OS=Aedes aegypti GN=PIWI7 PE=4 SV=1
CCG004026.1	4278	32	13	0.32141924	0.106231323	-1.597247078	Down	0.0004101	0.0006481	AAEL001543-PA OS=Aedes aegypti GN=AAEL001543 PE=4 SV=1
CCG000297.1	1125	32	13	1.22225023	0.403962309	-1.597247078	Down	0.0004101	0.00064802	AAEL009865-PA OS=Aedes aegypti GN=AAEL009865 PE=4 SV=1
CCG002856.1	1387	32	13	0.99137095	0.327655081	-1.597247078	Down	0.0004101	0.00064806	-
CCG023345.1	663	34	13	2.20357613	0.685456407	-1.684709919	Down	0.000153	0.0002474	AAEL002255-PA OS=Aedes aegypti GN=AAEL002255 PE=4 SV=1

CCG008316.1	1146	34	13	1.27484378	0.396559859	-1.684709919	Down	0.000153	0.00024738	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014452 PE=4 SV=1
CCG023322.1	492	34	13	2.9694532	0.923694305	-1.684709919	Down	0.000153	0.00024736	AAEL005184-PA OS=Aedes aegypti GN=AAEL005184 PE=4 SV=1
CCG016436.1	1515	35	13	0.99270014	0.299972012	-1.726530095	Down	9.25E-05	0.00015241	AAEL003653-PA OS=Aedes aegypti GN=SRPN12 PE=4 SV=2
CCG013982.1	1302	36	13	1.18810326	0.349045774	-1.767172079	Down	5.56E-05	9.26E-05	AAEL004231-PA OS=Aedes aegypti GN=AAEL004231 PE=4 SV=1
CCG016634.1	624	39	13	2.68560841	0.728297433	-1.882649297	Down	1.16E-05	2.02E-05	AAEL009567-PA OS=Aedes aegypti GN=AAEL009567 PE=4 SV=1
CCG001379.1	2847	42	13	0.63390546	0.159626835	-1.989564501	Down	2.30E-06	4.16E-06	AAEL005045-PA OS=Aedes aegypti GN=AAEL005045 PE=3 SV=1
CCG010918.1	723	43	13	2.55559971	0.628572058	-2.023511833	Down	1.33E-06	2.43E-06	AAEL000336-PA OS=Aedes aegypti GN=AAEL000336 PE=4 SV=1
CCG026239.1	1107	53	13	2.05726823	0.410530802	-2.325167533	Down	4.32E-09	8.72E-09	AAEL002954-PA OS=Aedes aegypti GN=AAEL002954 PE=4 SV=1
CCG018187.1	222	61	13	11.8069991	2.047106297	-2.527984416	Down	3.44E-11	7.45E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004854 PE=4 SV=1
CCG012482.1	576	78	13	5.81881823	0.788988885	-2.882649297	Down	7.02E-16	2.56E-15	-
CCG001823.1	2223	84	13	1.62368768	0.204434367	-2.989564501	Down	1.36E-17	5.19E-17	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013073 PE=4 SV=1
CCG001000.1	2333	87	13	1.60238616	0.19479537	-3.040190574	Down	1.86E-18	7.14E-18	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004811 PE=4 SV=1
CCG023396.1	1668	93	13	2.39579456	0.272456594	-3.136405889	Down	3.34E-20	1.31E-19	AAEL003564-PA OS=Aedes aegypti GN=AAEL003564 PE=4 SV=1
CCG015019.1	582	108	13	7.97376519	0.780854979	-3.35213458	Down	1.19E-24	4.84E-24	AAEL010664-PA OS=Aedes aegypti GN=AAEL010664 PE=3 SV=1
CCG003992.1	771	125	13	6.96655879	0.589439167	-3.563031363	Down	8.42E-30	3.54E-29	AAEL011457-PA OS=Aedes aegypti GN=AAEL011457 PE=4 SV=1

CCG005890.1	1212	183	13	6.48800448	0.374965015	-4.112946916	Down	5.82E-48	2.69E-47	AAEL012096-PA OS=Aedes aegypti GN=AAEL012096 PE=4 SV=1
CCG016355.1	198	279	13	60.5482624	2.295240394	-4.72136839	Down	4.31E-79	2.18E-78	-
CCG007166.1	417	281	13	28.9556245	1.089826374	-4.731673398	Down	9.49E-80	4.81E-79	AAEL004339-PA OS=Aedes aegypti GN=AAEL004339 PE=4 SV=1
CCG000876.1	1623	354	13	9.37232659	0.280010843	-5.064852628	Down	6.77E-104	3.61E-103	AAEL006339-PA OS=Aedes aegypti GN=AAEL006339 PE=4 SV=1
CCG026140.1	696	512	13	31.6099197	0.652956319	-5.597247078	Down	7.33E-157	4.20E-156	AAEL003501-PA OS=Aedes aegypti GN=AAEL003501 PE=4 SV=1
CCG021228.1	13725	929	13	2.90847967	0.033111665	-6.456781864	Down	1.02E-298	6.44E-298	AAEL007225-PA OS=Aedes aegypti GN=AAEL007225 PE=4 SV=1
CCG004326.1	1458	1098	13	32.3599236	0.311699313	-6.697909417	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000931 PE=4 SV=1
CCG016569.1	801	1202	13	64.4814245	0.567362794	-6.828468259	Down	0	0	AAEL010186-PA OS=Aedes aegypti GN=AAEL010186 PE=4 SV=1
CCG020646.1	271	0	14	0.001	1.805962638	10.81855233	Up	0.0002649	0.00042314	-
CCG015948.1	282	0	14	0.001	1.735517287	10.76115002	Up	0.0002649	0.00042325	AAEL005702-PA OS=Aedes aegypti GN=AAEL005702 PE=4 SV=1
CCG023240.1	309	0	14	0.001	1.583870145	10.62923834	Up	0.0002649	0.00042309	AAEL012826-PA OS=Aedes aegypti GN=AAEL012826 PE=4 SV=1
CCG001883.1	342	0	14	0.001	1.43104057	10.48284886	Up	0.0002649	0.00042344	-
CCG027975.1	351	0	14	0.001	1.394347222	10.44537415	Up	0.0002649	0.00042338	-
CCG014996.1	357	0	14	0.001	1.370912815	10.42092111	Up	0.0002649	0.0004229	-
CCG028611.1	528	0	14	0.001	0.926924005	9.856307253	Up	0.0002649	0.00042341	AAEL011712-PA OS=Aedes aegypti GN=AAEL011712 PE=4 SV=1
CCG004271.1	670	0	14	0.001	0.730471455	9.512684087	Up	0.0002649	0.00042336	Gag polyprotein OS=Anopheles gambiae PE=4 SV=1
CCG009507.1	732	0	14	0.001	0.668600922	9.385001534	Up	0.0002649	0.00042333	AAEL014647-PA OS=Aedes aegypti GN=AAEL014647 PE=4 SV=1
CCG028209.1	776	0	14	0.001	0.63069056	9.30078853	Up	0.0002649	0.00042293	AAEL009165-PA OS=Aedes aegypti GN=AAEL009165 PE=4 SV=1
CCG000163.1	788	0	14	0.001	0.621086136	9.278649553	Up	0.0002649	0.00042328	AAEL009165-PA OS=Aedes aegypti GN=AAEL009165 PE=4 SV=1

CCG000733.1	794	0	14	0.001	0.616392789	9.267706175	Up	0.0002649	0.00042312	AAEL013652-PA OS=Aedes aegypti GN=AAEL013652 PE=3 SV=1
CCG028103.1	816	0	14	0.001	0.599774356	9.22827603	Up	0.0002649	0.00042317	AAEL005607-PA OS=Aedes aegypti GN=AAEL005607 PE=3 SV=1
CCG012066.1	825	0	14	0.001	0.593231363	9.212451063	Up	0.0002649	0.00042306	AAEL012564-PA OS=Aedes aegypti GN=AAEL012564 PE=3 SV=1
CCG004721.2	832	0	14	0.001	0.588240234	9.200261654	Up	0.0002649	0.00042322	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014613 PE=4 SV=1
CCG022315.1	981	0	14	0.001	0.498894877	8.962592046	Up	0.0002649	0.00042301	Uncharacterized protein OS=Anopheles darlingi GN=AND_21982 PE=4 SV=1
CCG021767.1	1050	0	14	0.001	0.466110357	8.86452776	Up	0.0002649	0.00042298	Putative uncharacterized protein (Fragment) OS=Camponotus floridanus GN=EAG_00896 PE=4 SV=1
CCG003429.1	1257	0	14	0.001	0.389352327	8.604932438	Up	0.0002649	0.0004233	AAEL009473-PA OS=Aedes aegypti GN=AAEL009473 PE=4 SV=1
CCG021055.1	1398	0	14	0.001	0.350082886	8.451552727	Up	0.0002649	0.00042295	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_08164 PE=4 SV=1
CCG027428.1	1491	0	14	0.001	0.32824673	8.35863683	Up	0.0002649	0.0004232	AAEL005151-PA OS=Aedes aegypti GN=AAEL005151 PE=4 SV=1
CCG023244.1	1533	0	14	0.001	0.319253669	8.318559391	Up	0.0002649	0.00042304	AAEL007377-PA OS=Aedes aegypti GN=AAEL007377 PE=4 SV=1
CCG023757.1	878	34	14	1.66397606	0.55742127	-1.577794715	Down	0.0003044	0.00048409	AAEL002834-PA OS=Aedes aegypti GN=AAEL002834 PE=4 SV=1
CCG015189.1	696	34	14	2.09909623	0.703183728	-1.577794715	Down	0.0003044	0.00048406	Polyprotein-like OS=Solanum chilense PE=4 SV=1
CCG015091.1	459	34	14	3.1829433	1.066265522	-1.577794715	Down	0.0003044	0.00048412	AGAP003568-PA OS=Anopheles gambiae GN=AgaP_AGAP003568 PE=4 SV=4
CCG007135.1	960	36	14	1.61136505	0.509808203	-1.660256876	Down	0.0001148	0.00018677	AAEL000266-PA (Fragment) OS=Aedes aegypti GN=GPRNNA21 PE=4 SV=1

CCG002782.1	844	36	14	1.83283228	0.579876629	-1.660256876	Down	0.0001148	0.00018678	-
CCG011668.1	1068	36	14	1.44841802	0.458254564	-1.660256876	Down	0.0001148	0.00018676	AAEL004360-PA OS=Aedes aegypti GN=AAEL004360 PE=4 SV=1
CCG006193.1	1695	38	14	0.96333328	0.288740929	-1.738259388	Down	4.21E-05	7.10E-05	AAEL002067-PA (Fragment) OS=Aedes aegypti GN=AAEL002067 PE=3 SV=1
CCG000640.1	1864	38	14	0.87599244	0.262562165	-1.738259388	Down	4.21E-05	7.10E-05	AAEL009100-PA OS=Aedes aegypti GN=AAEL009100 PE=3 SV=1
CCG024318.1	714	38	14	2.28690464	0.685456407	-1.738259388	Down	4.21E-05	7.10E-05	AGAP013309-PA OS=Anopheles gambiae GN=AgaP_AGAP013309 PE=4 SV=1
CCG011465.1	627	39	14	2.67275861	0.780567583	-1.775734093	Down	2.53E-05	4.33E-05	AAEL009950-PA OS=Aedes aegypti GN=AAEL009950 PE=3 SV=1
CCG021832.1	627	40	14	2.74129088	0.780567583	-1.812259969	Down	1.51E-05	2.62E-05	AAEL009784-PA OS=Aedes aegypti GN=AAEL009784 PE=4 SV=1
CCG009935.1	2331	41	14	0.75579542	0.20995962	-1.847883879	Down	8.96E-06	1.57E-05	AAEL002732-PA (Fragment) OS=Aedes aegypti GN=AAEL002732 PE=4 SV=1
CCG027048.1	1350	44	14	1.40049505	0.362530278	-1.949763493	Down	1.81E-06	3.30E-06	AAEL005927-PA OS=Aedes aegypti GN=AAEL005927 PE=4 SV=1
CCG025631.1	1527	47	14	1.3225786	0.320508104	-2.044920726	Down	3.51E-07	6.59E-07	AAEL005090-PA OS=Aedes aegypti GN=CYP325X2 PE=3 SV=2
CCG014158.1	1548	48	14	1.33239487	0.316160126	-2.075294375	Down	2.02E-07	3.82E-07	AAEL001568-PA OS=Aedes aegypti GN=AAEL001568 PE=3 SV=1
CCG000793.1	228	55	14	10.3655062	2.146560854	-2.271691588	Down	3.68E-09	7.46E-09	AAEL012353-PA OS=Aedes aegypti GN=CTL15 PE=4 SV=1
CCG006150.1	1867	57	14	1.31187728	0.262140265	-2.323221888	Down	1.14E-09	2.34E-09	AAEL007619-PA OS=Aedes aegypti GN=TOLL5A PE=4 SV=1
CCG002262.1	489	60	14	5.27236007	1.00085046	-2.39722247	Down	1.90E-10	4.03E-10	AAEL009251-PA OS=Aedes aegypti GN=AAEL009251 PE=4 SV=1
CCG018068.1	1038	63	14	2.60798967	0.471498916	-2.467611798	Down	3.11E-11	6.76E-11	AAEL004106-PA OS=Aedes aegypti GN=AAEL004106 PE=4 SV=1
CCG008829.1	669	66	14	4.23916664	0.731563341	-2.534725993	Down	4.95E-12	1.13E-11	AAEL012952-PA OS=Aedes aegypti GN=AAEL012952 PE=1 SV=1

CCG019060.1	702	68	14	4.16231048	0.697173611	-2.577794715	Down	1.44E-12	3.48E-12	AAEL001208-PA OS=Aedes aegypti GN=AAEL001208 PE=4 SV=1
CCG001590.1	1578	68	14	1.85167424	0.310149477	-2.577794715	Down	1.44E-12	3.48E-12	AAEL009237-PA OS=Aedes aegypti GN=AAEL009237 PE=3 SV=1
CCG008493.1	621	75	14	5.18958147	0.788109299	-2.719150565	Down	1.77E-14	6.21E-14	AGAP011667-PA (Fragment) OS=Anopheles gambiae GN=AGAP011667 PE=4 SV=4
CCG018056.1	354	90	14	10.9245088	1.38253072	-2.98218497	Down	1.05E-18	4.03E-18	AAEL006695-PA OS=Aedes aegypti GN=AAEL006695 PE=4 SV=1
CCG001652.1	1323	100	14	3.24790133	0.369928855	-3.134188064	Down	1.32E-21	5.23E-21	AAEL007027-PA OS=Aedes aegypti GN=AAEL007027 PE=3 SV=1
CCG022077.1	780	102	14	5.61911914	0.62745625	-3.162757216	Down	3.43E-22	1.36E-21	Uncharacterized protein OS=Anopheles darlingi GN=AND_16384 PE=4 SV=1
CCG007766.1	1986	103	14	2.22854112	0.246432968	-3.176832401	Down	1.74E-22	6.95E-22	AAEL013732-PA (Fragment) OS=Aedes aegypti GN=AAEL013732 PE=4 SV=1
CCG023980.1	1401	110	14	3.37378359	0.349333244	-3.271691588	Down	1.49E-24	6.01E-24	Transporter OS=Aedes aegypti GN=AAEL000298 PE=3 SV=1
CCG024004.1	633	123	14	8.34956928	0.773168839	-3.432846379	Down	1.88E-28	7.86E-28	Lysosomal acid phosphatase OS=Culex quinquefasciatus GN=CpipJ_CPIJ008635 PE=4 SV=1
CCG015178.1	342	138	14	17.3386648	1.43104057	-3.598856331	Down	5.07E-33	2.17E-32	AAEL003048-PA OS=Aedes aegypti GN=AAEL003048 PE=4 SV=1
CCG015928.1	789	147	14	8.00576805	0.620298954	-3.690004219	Down	8.56E-36	3.73E-35	AAEL010639-PA OS=Aedes aegypti GN=AAEL010639 PE=4 SV=1
CCG005058.1	243	166	14	29.3538105	2.014057098	-3.865371305	Down	1.04E-41	4.65E-41	-
CCG024233.1	798	187	14	10.0693488	0.613303101	-4.037226334	Down	2.47E-48	1.14E-47	AAEL010639-PA OS=Aedes aegypti GN=AAEL010639 PE=4 SV=1
CCG010216.1	2568	225	14	3.76487161	0.190582506	-4.304113065	Down	1.73E-60	8.36E-60	AAEL011864-PA OS=Aedes aegypti GN=AAEL011864 PE=4 SV=1
CCG003871.1	908	249	14	11.7835506	0.539004267	-4.450333806	Down	2.99E-68	1.48E-67	Pyrroline-5-carboxylate reductase OS=Aedes aegypti PE=2 SV=1

CCG026910.1	1584	333	14	9.03341012	0.308974668	-4.869710241	Down	8.78E-96	4.61E-95	Glutamate dehydrogenase OS=Aedes aegypti GN=AAEL003757 PE=3 SV=1
CCG028322.1	1218	374	14	13.1943192	0.401819273	-5.037226334	Down	2.28E-109	1.22E-108	AAEL000044-PA OS=Aedes aegypti GN=CYP325Q1 PE=3 SV=1
CCG007954.1	1489	407	14	11.7452532	0.328687626	-5.159216858	Down	2.35E-120	1.29E-119	AAEL000729-PA OS=Aedes aegypti GN=AAEL005729 PE=4 SV=1
CCG010211.1	1239	1913	14	66.3447153	0.395008777	-7.391953032	Down	0	0	Vitelline envelope protein alpha OS=Oncorhynchus mykiss PE=2 SV=1
CCG008387.1	810	5349	14	283.759396	0.604217129	-8.875385362	Down	0	0	AAEL014369-PA OS=Aedes aegypti GN=AAEL014369 PE=4 SV=1
CCG001381.1	831	39144	14	2024.07616	0.588948104	-11.74683544	Down	0	0	-
CCG004491.1	362	0	15	0.001	1.44854738	10.50039116	Up	0.0001461	0.00023645	-
CCG018521.1	480	0	15	0.001	1.092446149	10.09334645	Up	0.0001461	0.00023639	AGAP013413-PA OS=Anopheles gambiae GN=AgaP_AGAP013413 PE=4 SV=1
CCG014603.1	504	0	15	0.001	1.040424904	10.02295712	Up	0.0001461	0.00023644	AAEL011252-PA OS=Aedes aegypti GN=AAEL011252 PE=4 SV=1
CCG021703.1	654	0	15	0.001	0.801795339	9.64709022	Up	0.0001461	0.00023642	AAEL010547-PA OS=Aedes aegypti GN=AAEL010547 PE=4 SV=1
CCG023512.1	756	0	15	0.001	0.693616603	9.437994622	Up	0.0001461	0.00023641	AAEL000465-PA OS=Aedes aegypti GN=AAEL005465 PE=4 SV=1
CCG028277.1	834	0	15	0.001	0.628745985	9.296333472	Up	0.0001461	0.0002365	AAEL000378-PA OS=Aedes aegypti GN=AAEL006378 PE=3 SV=1
CCG004308.1	951	0	15	0.001	0.551392378	9.106935515	Up	0.0001461	0.00023652	AAEL010140-PA OS=Aedes aegypti GN=AAEL010140 PE=3 SV=1
CCG011758.1	1188	0	15	0.001	0.441392383	8.785917925	Up	0.0001461	0.00023649	Odorant receptor 2 OS=Aedes albopictus PE=2 SV=1
CCG027607.1	1902	0	15	0.001	0.275696189	8.106935515	Up	0.0001461	0.00023647	Threonine--tRNA ligase OS=Wolbachia pipientis wAlbB GN=thrS PE=3 SV=1
CCG002187.1	300	34	15	4.86990326	1.747913839	-1.478259042	Down	0.0005769	0.00090336	-

CCG027298.1	675	35	15	2.22806031	0.776850595	-1.520079218	Down	0.0003622	0.00057348	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002736 PE=4 SV=1
CCG008149.1	453	36	15	3.41481335	1.157558834	-1.560721202	Down	0.0002258	0.00036181	AAEL000481-PA OS=Aedes aegypti GN=AAEL006481 PE=4 SV=1
CCG021867.1	548	42	15	3.29330083	0.956887138	-1.783113623	Down	1.15E-05	2.01E-05	-
CCG022228.1	588	43	15	3.14234454	0.891792775	-1.817060955	Down	6.89E-06	1.22E-05	LD21171p OS=Drosophila melanogaster PE=2 SV=1
CCG027456.1	432	45	15	4.47601402	1.213829055	-1.882649297	Down	2.42E-06	4.36E-06	ATPase 1120 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009335 PE=4 SV=1
CCG023947.1	847	45	15	2.28292569	0.619095811	-1.882649297	Down	2.42E-06	4.36E-06	-
CCG005919.1	1032	45	15	1.87368029	0.508114488	-1.882649297	Down	2.42E-06	4.36E-06	AAEL011623-PA OS=Aedes aegypti GN=AAEL011623 PE=4 SV=1
CCG010919.1	1113	51	15	1.96896358	0.471135806	-2.063221543	Down	9.31E-08	1.79E-07	AAEL000336-PA OS=Aedes aegypti GN=AAEL000336 PE=4 SV=1
CCG013838.1	1431	53	15	1.59147165	0.36643896	-2.118716655	Down	3.04E-08	5.97E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001348 PE=4 SV=1
CCG010740.1	744	57	15	3.29203612	0.704803967	-2.223686215	Down	3.10E-09	6.32E-09	AAEL013325-PA OS=Aedes aegypti GN=AAEL013325 PE=4 SV=1
CCG009366.1	710	60	15	3.63124518	0.738555143	-2.297686796	Down	5.41E-10	1.13E-09	-
CCG023860.1	729	64	15	3.77237725	0.719306106	-2.390796201	Down	5.04E-11	1.09E-10	AAEL013632-PA OS=Aedes aegypti GN=ABDA_AEDAE_b PE=2 SV=2
CCG027368.1	6264	65	15	0.44588645	0.083712349	-2.413164014	Down	2.77E-11	6.03E-11	Vitellogenin-C OS=Aedes aegypti GN=Vg-C PE=4 SV=1
CCG023579.1	948	73	15	3.30885087	0.553137291	-2.580620759	Down	2.08E-13	5.97E-13	AAEL011633-PA OS=Aedes aegypti GN=AAEL011633 PE=4 SV=1
CCG022667.1	1704	74	15	1.86605655	0.30773131	-2.600249566	Down	1.12E-13	3.47E-13	AAEL008532-PA (Fragment) OS=Aedes aegypti GN=AAEL008532 PE=4 SV=1
CCG000296.1	1314	82	15	2.68152073	0.399067086	-2.748348205	Down	7.18E-16	2.62E-15	AAEL001168-PA OS=Aedes aegypti GN=AAEL001168 PE=4 SV=1
CCG018916.1	859	82	15	4.10188386	0.610447208	-2.748348205	Down	7.18E-16	2.62E-15	-

CCG009809.1	447	82	15	7.88259114	1.173096536	-2.748348205	Down	7.18E-16	2.62E-15	AAEL005440-PA OS=Aedes aegypti GN=AAEL005440 PE=4 SV=1
CCG015018.1	558	90	15	6.93060236	0.939738623	-2.882649297	Down	4.12E-18	1.57E-17	AAEL010660-PA OS=Aedes aegypti GN=AAEL010660 PE=3 SV=1
CCG015833.1	2679	111	15	1.78038094	0.195735032	-3.185212067	Down	3.52E-24	1.42E-23	AAEL005810-PA OS=Aedes aegypti GN=AAEL005810 PE=4 SV=1
CCG012853.1	1626	127	15	3.35618468	0.322493328	-3.379480887	Down	6.12E-29	2.57E-28	AAEL003426-PA (Fragment) OS=Aedes aegypti GN=AAEL003426 PE=4 SV=1
CCG010572.1	558	134	15	10.3188968	0.939738623	-3.456885391	Down	4.73E-31	2.01E-30	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002735 PE=4 SV=1
CCG010860.2	1959	267	15	5.85651819	0.267674401	-4.451492132	Down	4.34E-73	2.17E-72	Hand domain-containing protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ015374 PE=4 SV=1
CCG018066.1	1494	287	15	8.2545608	0.350986715	-4.555703127	Down	1.34E-79	6.82E-79	AAEL004114-PA OS=Aedes aegypti GN=AAEL004114 PE=4 SV=1
CCG020462.1	1503	358	15	10.2349734	0.348884998	-4.874611978	Down	6.42E-103	3.41E-102	AAEL011445-PA OS=Aedes aegypti GN=AAEL011445 PE=4 SV=1
CCG020755.1	1908	1483	15	33.3983839	0.27482922	-6.925099083	Down	0	0	F-box/in protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011367 PE=4 SV=1
CCG018745.1	291	0	16	0.001	1.922104565	10.90847111	Up	8.05E-05	0.00013298	AAEL001787-PA OS=Aedes aegypti GN=AAEL001787 PE=4 SV=1
CCG019640.1	315	0	16	0.001	1.775658503	10.79413843	Up	8.05E-05	0.00013301	AAEL006670-PA OS=Aedes aegypti GN=AAEL006670 PE=4 SV=1
CCG020221.1	345	0	16	0.001	1.621253416	10.6628939	Up	8.05E-05	0.00013308	-
CCG009949.1	381	0	16	0.001	1.468064116	10.51969926	Up	8.05E-05	0.00013293	AAEL000073-PA OS=Aedes aegypti GN=AAEL000073 PE=4 SV=1
CCG023219.1	388	0	16	0.001	1.441578424	10.49343361	Up	8.05E-05	0.00013302	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG001247.1	390	0	16	0.001	1.434185714	10.48601614	Up	8.05E-05	0.00013296	-
CCG014442.1	507	0	16	0.001	1.10321978	10.10750451	Up	8.05E-05	0.00013306	AGAP004106-PA OS=Anopheles gambiae GN=AgaP_AGAP004106 PE=4 SV=5

CCG018342.1	513	0	16	0.001	1.090316624	10.09053143	Up	8.05E-05	0.00013307	-
CCG002251.1	576	0	16	0.001	0.971063244	9.923421449	Up	8.05E-05	0.00013299	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 SV=1
CCG012832.1	648	0	16	0.001	0.863167328	9.753496447	Up	8.05E-05	0.00013294	AAEL003547-PA OS=Aedes aegypti GN=AAEL003547 PE=4 SV=1
CCG021779.1	668	0	16	0.001	0.837323995	9.709642158	Up	8.05E-05	0.00013297	-
CCG022897.2	688	0	16	0.001	0.812983181	9.667081695	Up	8.05E-05	0.000133	-
CCG028412.1	696	0	16	0.001	0.803638546	9.650402954	Up	8.05E-05	0.00013307	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000496 PE=4 SV=1
CCG005480.2	819	0	16	0.001	0.682945578	9.415626809	Up	8.05E-05	0.000133	AAEL007279-PA OS=Aedes aegypti GN=AAEL007279 PE=3 SV=1
CCG022299.1	994	0	16	0.001	0.56270868	9.136244409	Up	8.05E-05	0.00013309	Putative uncharacterized protein (Fragment) OS=Harpegnathos saltator GN=EAI_06049 PE=4 SV=1
CCG022378.1	1116	0	16	0.001	0.501193932	8.969225138	Up	8.05E-05	0.00013303	AAEL008380-PA OS=Aedes aegypti GN=AAEL008380 PE=4 SV=1
CCG000478.1	1134	0	16	0.001	0.493238473	8.946141525	Up	8.05E-05	0.00013295	AAEL000130-PA OS=Aedes aegypti GN=AAEL000130 PE=4 SV=1
CCG003766.1	1254	0	16	0.001	0.446038619	8.801024817	Up	8.05E-05	0.00013294	AAEL001160-PA OS=Aedes aegypti GN=AAEL001160 PE=4 SV=1
CCG005674.1	1344	0	16	0.001	0.416169962	8.701029027	Up	8.05E-05	0.00013304	-
CCG014381.1	1608	0	16	0.001	0.34784355	8.442294759	Up	8.05E-05	0.00013305	AAEL006075-PA OS=Aedes aegypti GN=CYP307B1_a PE=3 SV=1
CCG004406.1	744	36	16	2.07918071	0.751790898	-1.467611798	Down	0.0004247	0.00067042	Putative uncharacterized protein (Fragment) OS=Camponotus floridanus GN=EAG_00524 PE=4 SV=1
CCG014901.1	1497	36	16	1.03334031	0.373635557	-1.467611798	Down	0.0004247	0.00067047	AAEL012766-PA OS=Aedes aegypti GN=CYP325G2 PE=3 SV=1
CCG023690.1	963	37	16	1.65096592	0.580822875	-1.507140162	Down	0.0002676	0.00042707	AAEL014518-PA OS=Aedes aegypti GN=AAEL014518 PE=4 SV=1
CCG005358.1	576	38	16	2.83480888	0.971063244	-1.54561431	Down	0.0001674	0.00026994	AAEL010802-PA OS=Aedes aegypti GN=AAEL010802 PE=4 SV=1

CCG008828.1	1018	40	16	1.68839822	0.549442464	-1.619614891	Down	6.43E-05	0.00010676	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017929 PE=4 SV=1
CCG012684.1	1596	40	16	1.0769357	0.350458915	-1.619614891	Down	6.43E-05	0.00010675	AAEL012356-PA (Fragment) OS=Aedes aegypti GN=GPRSMS PE=3 SV=2
CCG001989.1	1302	41	16	1.3531176	0.429594799	-1.655238801	Down	3.94E-05	6.66E-05	AAEL003209-PA OS=Aedes aegypti GN=AAEL003209 PE=4 SV=1
CCG014253.1	2556	43	16	0.72288677	0.218831154	-1.723951551	Down	1.46E-05	2.53E-05	AAEL008549-PA OS=Aedes aegypti GN=AAEL008549 PE=4 SV=1
CCG021470.1	1337	44	16	1.41411243	0.418348862	-1.757118415	Down	8.80E-06	1.54E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_11969 PE=4 SV=1
CCG024686.1	732	44	16	2.58288022	0.764115339	-1.757118415	Down	8.80E-06	1.54E-05	AAEL011575-PA OS=Aedes aegypti GN=AAEL011575 PE=4 SV=1
CCG003724.1	1254	49	16	1.67904067	0.446038619	-1.91239664	Down	6.51E-07	1.21E-06	AAEL007607-PA OS=Aedes aegypti GN=CYP4D38 PE=3 SV=1
CCG011890.1	621	50	16	3.45972098	0.900696342	-1.941542986	Down	3.81E-07	7.14E-07	AAEL000340-PA OS=Aedes aegypti GN=AAEL000340 PE=3 SV=1
CCG021220.1	816	52	16	2.7382674	0.685456407	-1.998126514	Down	1.29E-07	2.47E-07	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004860 PE=3 SV=1
CCG011224.1	393	52	16	5.68556285	1.423237731	-1.998126514	Down	1.29E-07	2.47E-07	-
CCG021602.1	186	56	16	12.9371244	3.007163593	-2.105041718	Down	1.41E-08	2.80E-08	AAEL009548-PA OS=Aedes aegypti GN=AAEL009548 PE=4 SV=1
CCG026953.1	735	68	16	3.97543123	0.760996501	-2.385149637	Down	1.34E-11	2.96E-11	AAEL005072-PA OS=Aedes aegypti GN=AAEL005072 PE=4 SV=1
CCG009992.1	1905	69	16	1.55638409	0.293612823	-2.406211253	Down	7.35E-12	1.65E-11	AAEL012949-PA (Fragment) OS=Aedes aegypti GN=AAEL012949 PE=4 SV=1
CCG002183.1	1506	72	16	2.05432994	0.371402675	-2.467611798	Down	1.20E-12	2.93E-12	AAEL007673-PA OS=Aedes aegypti GN=AAEL007673 PE=3 SV=1
CCG001553.1	1458	76	16	2.23984899	0.383629923	-2.54561431	Down	1.04E-13	3.25E-13	AAEL001902-PA OS=Aedes aegypti GN=AAEL001902 PE=3 SV=1

CCG014783.1	519	82	16	6.78905248	1.077711808	-2.655238801	Down	2.48E-15	8.97E-15	AAEL009978-PA OS=Aedes aegypti GN=AAEL009978 PE=4 SV=1
CCG005496.1	1452	88	16	2.60422634	0.385215171	-2.757118415	Down	5.53E-17	2.09E-16	AAEL008053-PA OS=Aedes aegypti GN=AAEL008053 PE=3 SV=1
CCG003240.1	1908	93	16	2.09443675	0.293151168	-2.836845607	Down	2.22E-18	8.51E-18	AAEL010851-PA OS=Aedes aegypti GN=AAEL010851 PE=4 SV=1
CCG008619.1	741	96	16	5.56692918	0.754834586	-2.882649297	Down	3.17E-19	1.23E-18	AGAP011666-PA (Fragment) OS=Anopheles gambiae GN=AGAP011666 PE=4 SV=2
CCG007376.1	985	104	16	4.53690599	0.567850181	-2.998126514	Down	1.66E-21	6.55E-21	AAEL011634-PA OS=Aedes aegypti GN=AAEL011634 PE=4 SV=1
CCG009288.1	900	105	16	5.0131357	0.621480476	-3.011932314	Down	8.54E-22	3.39E-21	AAEL002673-PA OS=Aedes aegypti GN=AAEL002673 PE=4 SV=1
CCG015759.1	1305	105	16	3.45733497	0.428607225	-3.011932314	Down	8.54E-22	3.39E-21	AAEL007016-PA OS=Aedes aegypti GN=AAEL007016 PE=3 SV=1
CCG002341.1	2469	108	16	1.87959957	0.226542093	-3.052574298	Down	1.16E-22	4.64E-22	AAEL014202-PA OS=Aedes aegypti GN=AAEL014202 PE=4 SV=1
CCG009684.1	609	109	16	7.69080636	0.918444053	-3.065871121	Down	5.97E-23	2.39E-22	AAEL001563-PA OS=Aedes aegypti GN=AAEL001563 PE=4 SV=1
CCG003247.1	1911	110	16	2.47340178	0.292690962	-3.07904651	Down	3.06E-23	1.23E-22	AAEL013732-PA (Fragment) OS=Aedes aegypti GN=AAEL013732 PE=4 SV=1
CCG026968.1	1353	115	16	3.65226865	0.413401647	-3.143176847	Down	1.07E-24	4.33E-24	AAEL008726-PA OS=Aedes aegypti GN=AAEL008726 PE=4 SV=1
CCG003446.1	468	117	16	10.7424337	1.195154761	-3.168051516	Down	2.76E-25	1.13E-24	-
CCG008135.1	747	153	16	8.80102998	0.748771658	-3.555074639	Down	4.35E-36	1.90E-35	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017322 PE=4 SV=1
CCG014925.1	831	179	16	9.25581528	0.673083548	-3.781502573	Down	4.10E-44	1.86E-43	AAEL004082-PA OS=Aedes aegypti GN=AAEL004082 PE=3 SV=1
CCG002894.1	1839	184	16	4.29931004	0.304150314	-3.821248752	Down	1.13E-45	5.18E-45	AAEL011809-PA OS=Aedes aegypti GN=AAEL011809 PE=3 SV=1
CCG013257.1	669	188	16	12.075202	0.836072389	-3.852275648	Down	6.35E-47	2.92E-46	AAEL001233-PA OS=Aedes aegypti GN=CLIFE9 PE=3 SV=2
CCG027022.1	543	238	16	18.83388	1.030078137	-4.192504559	Down	9.07E-63	4.41E-62	-

CCG020424.1	1239	337	16	11.6874904	0.451438602	-4.694291577	Down	6.65E-95	3.49E-94	AAEL012794-PA OS=Aedes aegypti GN=AAEL012794 PE=4 SV=1
CCG005485.1	1422	607	16	18.3422144	0.393342073	-5.543239502	Down	6.62E-185	3.90E-184	AAEL005004-PA OS=Aedes aegypti GN=AAEL005004 PE=4 SV=1
CCG004395.1	228	0	17	0.001	2.60653818	11.34791928	Up	4.44E-05	7.47E-05	-
CCG007233.1	429	0	17	0.001	1.385293019	10.43597545	Up	4.44E-05	7.48E-05	-
CCG004633.1	442	0	17	0.001	1.344549107	10.39290673	Up	4.44E-05	7.47E-05	Putative secreted protein OS=Aedes aegypti PE=2 SV=1
CCG016358.1	627	0	17	0.001	0.947832066	9.888487659	Up	4.44E-05	7.48E-05	AAEL004696-PA OS=Aedes aegypti GN=AAEL004696 PE=4 SV=1
CCG009755.1	664	0	17	0.001	0.895016122	9.80576986	Up	4.44E-05	7.47E-05	-
CCG012375.1	737	0	17	0.001	0.806364593	9.655288482	Up	4.44E-05	7.47E-05	AAEL007959-PA OS=Aedes aegypti GN=AAEL007959 PE=4 SV=1
CCG020709.1	1017	0	17	0.001	0.584356642	9.190705328	Up	4.44E-05	7.47E-05	AAEL005443-PA (Fragment) OS=Aedes aegypti GN=AAEL005443 PE=4 SV=1
CCG015029.1	1044	0	17	0.001	0.56924397	9.152903295	Up	4.44E-05	7.47E-05	AAEL008380-PA OS=Aedes aegypti GN=AAEL008380 PE=4 SV=1
CCG001620.1	1065	0	17	0.001	0.558019441	9.124171576	Up	4.44E-05	7.47E-05	AAEL004590-PA OS=Aedes aegypti GN=AAEL004590 PE=4 SV=1
CCG017233.1	1170	0	17	0.001	0.507940774	8.988516477	Up	4.44E-05	7.47E-05	AAEL015312-PA OS=Aedes aegypti GN=AAEL015312 PE=3 SV=1
CCG009838.1	1179	0	17	0.001	0.504063363	8.977461288	Up	4.44E-05	7.47E-05	PA D-oxidoreductase OS=Culex quinquefasciatus GN=CpipJ_CPIJ016820 PE=4 SV=1
CCG022990.1	1267	0	17	0.001	0.469053437	8.873608482	Up	4.44E-05	7.47E-05	RNA nuclease-reverse transcriptase OS=Bombyx mori PE=4 SV=1
CCG010527.1	1308	0	17	0.001	0.454350692	8.827662466	Up	4.44E-05	7.48E-05	AGAP000437-PA OS=Anopheles gambiae GN=AgaP_AGAP000437 PE=4 SV=5
CCG006770.1	2919	0	17	0.001	0.203593938	7.669550796	Up	4.44E-05	7.47E-05	AAEL005942-PA OS=Aedes aegypti GN=AAEL005942 PE=4 SV=1
CCG027149.1	1134	38	17	1.43990292	0.524065878	-1.458151468	Down	0.0003128	0.00049712	Calvar serum protein 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009032 PE=4 SV=1

CCG000116.1	1356	38	17	1.2041666	0.438267482	-1.458151468	Down	0.0003128	0.00049709	AAEL006886-PA OS=Aedes aegypti GN=AAEL006886 PE=4 SV=1
CCG012645.1	843	40	17	2.03889607	0.70497118	-1.53215205	Down	0.0001241	0.00020171	AAEL013324-PA OS=Aedes aegypti GN=AAEL013324 PE=4 SV=1
CCG019643.1	1521	41	17	1.15829002	0.390723672	-1.56777596	Down	7.74E-05	0.00012806	AAEL009780-PA OS=Aedes aegypti GN=AAEL009780 PE=3 SV=1
CCG013704.1	798	41	17	2.20771819	0.744725194	-1.56777596	Down	7.74E-05	0.00012806	AAEL001809-PA OS=Aedes aegypti GN=AAEL001809 PE=4 SV=1
CCG005688.1	2316	41	17	0.76069047	0.256602204	-1.56777596	Down	7.74E-05	0.00012805	AAEL002066-PA OS=Aedes aegypti GN=AAEL002066 PE=4 SV=1
CCG018402.1	2469	42	17	0.73095539	0.240700974	-1.602541378	Down	4.80E-05	8.06E-05	AAEL002338-PA OS=Aedes aegypti GN=AAEL002338 PE=4 SV=1
CCG027164.1	1290	43	17	1.43232449	0.460690469	-1.63648871	Down	2.96E-05	5.03E-05	-
CCG008428.1	525	47	17	3.84681434	1.131982295	-1.764812807	Down	4.03E-06	7.22E-06	AAEL013104-PA OS=Aedes aegypti GN=AAEL013104 PE=4 SV=1
CCG006161.1	593	53	17	3.84046532	1.002176569	-1.938144409	Down	1.75E-07	3.33E-07	-
CCG007370.1	3549	55	17	0.66591586	0.167453002	-1.991583668	Down	5.96E-08	1.16E-07	AAEL007976-PA OS=Aedes aegypti GN=AAEL007976 PE=4 SV=1
CCG017262.1	342	56	17	7.03597994	1.73769212	-2.017578877	Down	3.45E-08	6.75E-08	AAEL008028-PA OS=Aedes aegypti GN=AAEL008028 PE=4 SV=1
CCG009275.1	863	72	17	3.58496048	0.688633494	-2.380148956	Down	3.55E-12	8.26E-12	AAEL015464-PA OS=Aedes aegypti GN=AAEL015464 PE=3 SV=1
CCG012013.1	1455	87	17	2.56932434	0.40844722	-2.653167451	Down	3.60E-16	1.35E-15	AAEL011160-PA OS=Aedes aegypti GN=AAEL011160 PE=4 SV=1
CCG003421.1	900	106	17	5.06087985	0.660323006	-2.938144409	Down	1.79E-21	7.05E-21	AAEL003279-PA OS=Aedes aegypti GN=AAEL003279 PE=3 SV=1
CCG007496.1	1782	112	17	2.70067917	0.333496468	-3.017578877	Down	3.41E-23	1.37E-22	AAEL015391-PA OS=Aedes aegypti GN=AAEL015391 PE=4 SV=1
CCG017844.1	1307	114	17	3.74793401	0.454698321	-3.043113969	Down	9.02E-24	3.63E-23	AAEL002869-PA OS=Aedes aegypti GN=AAEL002869 PE=4 SV=1
CCG024917.1	3018	117	17	1.6658247	0.196915409	-3.080588675	Down	1.22E-24	4.94E-24	AAEL006760-PA (Fragment) OS=Aedes aegypti GN=AAEL006760 PE=4 SV=1
CCG019175.1	499	118	17	10.1611797	1.190963337	-3.092867004	Down	6.25E-25	2.54E-24	AAEL002591-PA OS=Aedes aegypti GN=AAEL002591 PE=4 SV=1

CCG006301.1	1203	175	17	6.25079265	0.494007236	-3.661435067	Down	4.53E-42	2.04E-41	AAEL004711-PA OS=Aedes aegypti GN=AAEL004711 PE=4 SV=1
CCG005225.1	519	194	17	16.0619047	1.145068796	-3.810136797	Down	5.76E-48	2.66E-47	AAEL009978-PA OS=Aedes aegypti GN=AAEL009978 PE=4 SV=1
CCG015168.1	1029	218	17	9.10340345	0.577541987	-3.97840828	Down	1.67E-55	7.94E-55	AAEL003411-PA OS=Aedes aegypti GN=AAEL003411 PE=4 SV=1
CCG024377.1	600	278	17	19.9093104	0.990484509	-4.329165028	Down	1.13E-74	5.67E-74	AAEL009112-PA OS=Aedes aegypti GN=AAEL009112 PE=4 SV=1
CCG000946.1	771	330	17	18.3917152	0.770805065	-4.576546169	Down	1.49E-91	7.75E-91	AAEL008152-PA OS=Aedes aegypti GN=AAEL008152 PE=4 SV=1
CCG011695.1	954	405	17	18.2418685	0.622946232	-4.872002053	Down	3.39E-116	1.84E-115	AAEL013222-PA OS=Aedes aegypti GN=AAEL013222 PE=4 SV=1
CCG015449.1	1491	425	17	12.2482476	0.398585315	-4.941542986	Down	8.20E-123	4.51E-122	AAEL012701-PA OS=Aedes aegypti GN=CYP325T2 PE=3 SV=1
CCG015583.1	2148	538	17	10.7624382	0.27667165	-5.281686317	Down	1.89E-160	1.09E-159	AAEL006245-PA (Fragment) OS=Aedes aegypti GN=AAEL006245 PE=4 SV=1
CCG000282.1	756	2184	17	124.134789	0.786098816	-7.302981096	Down	0	0	AAEL011797-PA OS=Aedes aegypti GN=AAEL011797 PE=4 SV=1
CCG007816.1	711	4427	17	267.548544	0.835851906	-8.322337613	Down	0	0	AAEL009577-PA OS=Aedes aegypti GN=AAEL009577 PE=4 SV=1
CCG002351.1	306	6399	17	898.57298	1.942126487	-8.853854706	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015149 PE=4 SV=1
CCG004740.1	282	0	18	0.001	2.231379368	11.1237201	Up	2.45E-05	4.20E-05	AAEL000685-PA OS=Aedes aegypti GN=AAEL000685 PE=4 SV=1
CCG004140.1	528	0	18	0.001	1.191759435	10.21887733	Up	2.45E-05	4.20E-05	-
CCG027637.1	546	0	18	0.001	1.152470663	10.17051431	Up	2.45E-05	4.20E-05	AAEL011839-PA OS=Aedes aegypti GN=AAEL011839 PE=4 SV=1
CCG004677.1	690	0	18	0.001	0.911955046	9.8328189	Up	2.45E-05	4.20E-05	AAEL001814-PA OS=Aedes aegypti GN=AAEL001814 PE=4 SV=1
CCG015311.1	731	0	18	0.001	0.860805721	9.749543856	Up	2.45E-05	4.20E-05	Hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG001873.1	885	0	18	0.001	0.711015799	9.473737807	Up	2.45E-05	4.20E-05	AAEL008197-PA OS=Aedes aegypti GN=AAEL008197 PE=3 SV=1

CCG015533.1	909	0	18	0.001	0.692243104	9.435134967	Up	2.45E-05	4.20E-05	AAEL012550-PA OS=Aedes aegypti GN=AAEL012550 PE=4 SV=1
CCG003452.1	933	0	18	0.001	0.674436208	9.397538181	Up	2.45E-05	4.20E-05	AAEL000224-PA OS=Aedes aegypti GN=AAEL000224 PE=3 SV=1
CCG006009.1	1170	0	18	0.001	0.537819643	9.070978637	Up	2.45E-05	4.20E-05	3-demethylubiquinone-9 3-methyltransferase OS=Wolbachia pipientis wAlbB GN=ubiG PE=3 SV=1
CCG002815.1	1216	0	18	0.001	0.517474492	9.015343938	Up	2.45E-05	4.20E-05	-
CCG016924.1	1494	0	18	0.001	0.421184057	8.718307019	Up	2.45E-05	4.20E-05	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG023831.1	1686	0	18	0.001	0.373220037	8.543882631	Up	2.45E-05	4.20E-05	AAEL003201-PA (Fragment) OS=Aedes aegypti GN=AAEL003201 PE=4 SV=1
CCG011776.1	2614	0	18	0.001	0.24072264	7.911228026	Up	2.45E-05	4.20E-05	RNase H and integrase-like protein (Fragment) OS=Bombyx mori GN=P-1 PE=4 SV=1
CCG024250.1	342	41	18	5.15134245	1.839909304	-1.485313799	Down	0.0001461	0.00023641	AAEL003755-PA OS=Aedes aegypti GN=AAEL003755 PE=3 SV=1
CCG014953.1	1116	43	18	1.6556439	0.563843174	-1.554026549	Down	5.76E-05	9.59E-05	AAEL009865-PA OS=Aedes aegypti GN=AAEL009865 PE=4 SV=1
CCG010552.1	729	43	18	2.53456596	0.863167328	-1.554026549	Down	5.76E-05	9.59E-05	AGAP010049-PA (Fragment) OS=Anopheles gambiae GN=AGAP010049 PE=4 SV=4
CCG010104.1	468	45	18	4.13170525	1.344549107	-1.619614891	Down	2.22E-05	3.81E-05	-
CCG021530.1	921	46	18	2.14615395	0.68322365	-1.651323751	Down	1.36E-05	2.37E-05	AAEL012170-PA OS=Aedes aegypti GN=AAEL012170 PE=4 SV=1
CCG018165.1	408	48	18	5.0552629	1.542276916	-1.712724295	Down	5.07E-06	9.02E-06	Uncharacterized protein OS=Anopheles darlingi GN=AND_07352 PE=4 SV=1
CCG017544.1	507	49	18	4.15289348	1.241122252	-1.742471639	Down	3.07E-06	5.52E-06	AAEL014490-PA OS=Aedes aegypti GN=AAEL014490 PE=4 SV=1
CCG010306.1	2208	52	18	1.01196839	0.284985952	-1.828201513	Down	6.63E-07	1.23E-06	AAEL002676-PA OS=Aedes aegypti GN=AAEL002676 PE=4 SV=1
CCG007103.1	788	54	18	2.94462648	0.798539317	-1.882649297	Down	2.33E-07	4.41E-07	-

CCG014766.1	897	54	18	2.58680677	0.701503882	-1.882649297	Down	2.33E-07	4.41E-07	AAEL004569-PA OS=Aedes aegypti GN=AAEL004569 PE=4 SV=1
CCG013106.1	216	55	18	10.9413676	2.913189731	-1.909121508	Down	1.37E-07	2.62E-07	AAEL002011-PA OS=Aedes aegypti GN=AAEL002011 PE=4 SV=1
CCG028507.1	879	58	18	2.8353181	0.715869149	-1.98574279	Down	2.75E-08	5.40E-08	Pv-like protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011811 PE=4 SV=1
CCG021114.2	1453	64	18	1.89267929	0.433068811	-2.127761795	Down	9.96E-10	2.05E-09	Peptidase OS=Culex quinquefasciatus GN=CpipJ_CPIJ014636 PE=4 SV=1
CCG028474.1	1440	67	18	1.99928626	0.43697846	-2.193850985	Down	1.81E-10	3.84E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018073 PE=4 SV=1
CCG006885.1	396	75	18	8.13820731	1.589012581	-2.356580485	Down	1.71E-12	4.10E-12	AAEL001463-PA OS=Aedes aegypti GN=AAEL001463 PE=4 SV=1
CCG021344.1	306	84	18	11.7956134	2.056369222	-2.520079218	Down	7.50E-15	2.69E-14	Uncharacterized protein OS=Anopheles darlingi GN=AND_25223 PE=4 SV=1
CCG019322.1	432	90	18	8.95202804	1.456594866	-2.619614891	Down	1.83E-16	6.88E-16	AAEL014559-PA OS=Aedes aegypti GN=AAEL014559 PE=4 SV=1
CCG017170.1	1512	95	18	2.69981798	0.416169962	-2.697617403	Down	7.92E-18	3.02E-17	AAEL000912-PA OS=Aedes aegypti GN=AAEL000912 PE=4 SV=1
CCG014828.1	261	111	18	18.2744848	2.410915639	-2.922177661	Down	2.65E-22	1.06E-21	AAEL009481-PA OS=Aedes aegypti GN=AAEL009481 PE=4 SV=1
CCG012841.1	1035	122	18	5.06503152	0.607970031	-3.058499132	Down	1.85E-25	7.55E-25	AAEL013276-PA OS=Aedes aegypti GN=AAEL013276 PE=4 SV=1
CCG007131.1	2292	126	18	2.36221054	0.274541441	-3.105041718	Down	1.28E-26	5.26E-26	AAEL012309-PA OS=Aedes aegypti GN=AAEL012309 PE=4 SV=1
CCG011561.1	1725	133	18	3.31302881	0.364782018	-3.18304423	Down	1.14E-28	4.78E-28	AAEL004766-PA OS=Aedes aegypti GN=AAEL004766 PE=4 SV=1
CCG008132.1	1482	152	18	4.40715227	0.424594455	-3.375689308	Down	2.57E-34	1.11E-33	AAEL004213-PA OS=Aedes aegypti GN=AAEL004213 PE=4 SV=1
CCG015784.1	1569	180	18	4.92960627	0.401050976	-3.619614891	Down	7.92E-43	3.57E-42	AAEL007684-PA OS=Aedes aegypti GN=AAEL007684 PE=4 SV=1

CCG020031.1	1146	246	18	9.22386973	0.549082881	-4.0702763	Down	1.73E-63	8.42E-63	Protoninament ribbon protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001618 PE=4 SV=1
CCG000346.1	1248	357	18	12.2918231	0.504205915	-4.607542059	Down	2.50E-99	1.32E-98	AAEL005969-PA OS=Aedes aegypti GN=AAEL005969 PE=4 SV=1
CCG000947.1	771	458	18	25.5254714	0.816146539	-4.966965583	Down	1.35E-132	7.51E-132	AAEL008159-PA OS=Aedes aegypti GN=AAEL008159 PE=4 SV=1
CCG026890.1	1614	500	18	13.3115659	0.389869258	-5.093546079	Down	1.51E-146	8.52E-146	AAEL005533-PA (Fragment) OS=Aedes aegypti GN=AAEL005533 PE=4 SV=1
CCG018766.1	1797	935	18	22.3576527	0.350166378	-5.99658435	Down	3.53E-293	2.21E-292	AGAP005519-PA OS=Anopheles gambiae GN=AGAP005519 PE=4 SV=3
CCG000708.1	981	1963	18	85.9832712	0.641436271	-7.066606252	Down	0	0	AAEL000432-PA OS=Aedes aegypti GN=AAEL000432 PE=3 SV=1
CCG015395.1	624	11910	18	820.143492	1.00841183	-9.667647587	Down	0	0	AAEL004292-PA OS=Aedes aegypti GN=AAEL004292 PE=4 SV=1
CCG014096.1	273	0	19	0.001	2.432993622	11.24851682	Up	1.35E-05	2.35E-05	AAEL004628-PA OS=Aedes aegypti GN=AAEL004628 PE=4 SV=1
CCG017730.1	456	0	19	0.001	1.456594866	10.50838395	Up	1.35E-05	2.35E-05	AAEL009598-PA OS=Aedes aegypti GN=AAEL009598 PE=4 SV=1
CCG004069.1	573	0	19	0.001	1.159174972	10.17888263	Up	1.35E-05	2.35E-05	AAEL012461-PA OS=Aedes aegypti GN=AAEL012461 PE=4 SV=1
CCG017058.1	840	0	19	0.001	0.790722927	9.627028446	Up	1.35E-05	2.35E-05	AAEL008797-PA OS=Aedes aegypti GN=AAEL008797 PE=4 SV=1
CCG010721.1	978	0	19	0.001	0.679148526	9.407583309	Up	1.35E-05	2.35E-05	AAEL009463-PA (Fragment) OS=Aedes aegypti GN=AAEL009463 PE=4 SV=1
CCG003303.1	1032	0	19	0.001	0.643611685	9.330046708	Up	1.35E-05	2.35E-05	AAEL009977-PA OS=Aedes aegypti GN=AAEL009977 PE=4 SV=1
CCG008512.1	1215	0	19	0.001	0.546672641	9.094533365	Up	1.35E-05	2.35E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014988 PE=3 SV=1

CCG001859.1	1434	0	19	0.001	0.463184978	8.855444655	Up	1.35E-05	2.35E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017585 PE=4 SV=1
CCG010597.1	1438	39	19	1.16538223	0.461896564	-1.335161502	Down	0.0006364	0.00098882	AAEL0008020-FA OS=Aedes aegypti GN=AAEL008626 PE=4 SV=1
CCG024758.1	1725	39	19	0.97148965	0.385047686	-1.335161502	Down	0.0006364	0.00098888	AAEL0005255-FA OS=Aedes aegypti GN=AAEL003233 PE=3 SV=1
CCG002128.1	1096	40	19	1.56823849	0.606028521	-1.371687378	Down	0.0004126	0.00065184	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG027899.1	732	41	19	2.40677475	0.907386965	-1.407311287	Down	0.0002656	0.00042403	AAEL0005840-FA OS=Aedes aegypti GN=AAEL005840 PE=4 SV=1
CCG022498.1	648	41	19	2.71876407	1.025011202	-1.407311287	Down	0.0002656	0.00042406	AAEL0009546-FA OS=Aedes aegypti GN=AAEL009548 PE=4 SV=1
CCG019750.1	318	42	19	5.67524797	2.088702071	-1.442076706	Down	0.0001699	0.00027381	AAEL0009314-FA OS=Aedes aegypti GN=AAEL009314 PE=4 SV=1
CCG012309.2	1689	44	19	1.11940102	0.393254742	-1.509190901	Down	6.82E-05	0.00011315	GATA transcription factor GATAd OS=Aedes aegypti PE=2 SV=1
CCG012525.1	528	45	19	3.66219329	1.257968293	-1.541612379	Down	4.28E-05	7.21E-05	-
CCG000145.1	1239	47	19	1.63000607	0.53608334	-1.604348134	Down	1.66E-05	2.86E-05	AAEL0009966-FA OS=Aedes aegypti GN=AAEL009966 PE=4 SV=1
CCG001314.1	1548	48	19	1.33239487	0.429074457	-1.634721783	Down	1.02E-05	1.79E-05	AAEL000127-FA OS=Aedes aegypti GN=AAEL000127 PE=4 SV=1
CCG021910.1	1512	48	19	1.36411856	0.439290515	-1.634721783	Down	1.02E-05	1.79E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002477 PE=4 SV=1
CCG011892.1	984	55	19	2.40176362	0.675007377	-1.831118996	Down	3.05E-07	5.72E-07	AAEL000336-FA OS=Aedes aegypti GN=AAEL000336 PE=4 SV=1
CCG010724.2	1939	57	19	1.26316394	0.342551449	-1.882649297	Down	1.07E-07	2.06E-07	Protein translocase subunit secA OS=Harpegnathos saltator GN=EAI_00772 PE=4 SV=1
CCG022426.1	495	58	19	5.03483759	1.341832846	-1.907740278	Down	6.33E-08	1.23E-07	AAEL0006630-FA OS=Aedes aegypti GN=AAEL000630 PE=4 SV=1
CCG008044.1	3204	59	19	0.7912654	0.207305636	-1.932402332	Down	3.72E-08	7.26E-08	-

CCG024023.1	858	60	19	3.00487655	0.774134334	-1.956649878	Down	2.18E-08	4.29E-08	AGAP000352-PA OS=Anopheles gambiae GN=AgaP_AGAP000352 PE=4 SV=2
CCG006546.1	582	60	19	4.42986955	1.141249585	-1.956649878	Down	2.18E-08	4.29E-08	AAEL010664-PA OS=Aedes aegypti GN=AAEL010664 PE=3 SV=1
CCG007501.1	2357	62	19	1.13030273	0.281801977	-2.003955593	Down	7.38E-09	1.48E-08	Ankyrin repeat protein OS=Neosartorya fumigata (strain CEA10 / CBS 144.89 / FGSC A1163) GN=AFUB_080240 PE=4 SV=1
CCG021589.1	1500	64	19	1.83337534	0.442804839	-2.049759283	Down	2.47E-09	5.03E-09	AAEL008933-PA OS=Aedes aegypti GN=AAEL008933 PE=4 SV=1
CCG023587.1	789	66	19	3.59442647	0.841834295	-2.094153402	Down	8.13E-10	1.68E-09	AAEL009032-PA OS=Aedes aegypti GN=AAEL009032 PE=4 SV=1
CCG013492.1	543	69	19	5.46024252	1.223217788	-2.15828374	Down	1.51E-10	3.20E-10	AAEL010347-PA OS=Aedes aegypti GN=AAEL010347 PE=4 SV=1
CCG019533.1	1014	73	19	3.09348188	0.655036744	-2.239583842	Down	1.53E-11	3.37E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_17189 PE=4 SV=1
CCG010011.1	786	75	19	4.10016552	0.845047403	-2.278577973	Down	4.78E-12	1.09E-11	AAEL002684-PA OS=Aedes aegypti GN=AAEL002684 PE=4 SV=1
CCG004011.1	660	80	19	5.20845268	1.006374634	-2.371687378	Down	2.51E-13	7.06E-13	Putative reverse transcriptase OS=Anopheles gambiae PE=4 SV=1
CCG005422.1	1968	83	19	1.81223982	0.337503688	-2.424798714	Down	4.18E-14	1.42E-13	AAEL010471-PA OS=Aedes aegypti GN=AAEL010471 PE=3 SV=1
CCG003040.1	1083	86	19	3.41218576	0.613303101	-2.476024037	Down	6.82E-15	2.45E-14	AAEL013496-PA OS=Aedes aegypti GN=PPO8 PE=4 SV=1
CCG013568.1	381	87	19	9.81198664	1.743326138	-2.492702779	Down	3.71E-15	1.34E-14	AAEL002231-PA OS=Aedes aegypti GN=AAEL002231 PE=4 SV=1
CCG011780.1	2070	88	19	1.82673268	0.320873072	-2.509190901	Down	2.01E-15	7.30E-15	AAEL011764-PA OS=Aedes aegypti GN=PPO10 PE=4 SV=1
CCG027721.1	1384	96	19	2.98055963	0.47991854	-2.634721783	Down	1.43E-17	5.44E-17	-
CCG018922.1	2385	102	19	1.83769934	0.27849361	-2.722184625	Down	3.26E-19	1.26E-18	AAEL000912-PA OS=Aedes aegypti GN=AAEL000912 PE=4 SV=1

CCG009262.1	717	109	19	6.53235854	0.926369956	-2.817943608	Down	3.70E-21	1.46E-20	AAEL005165-PA OS=Aedes aegypti GN=AAEL003183 PE=3 SV=1
CCG015685.1	1074	110	19	4.40099703	0.618442513	-2.831118996	Down	1.94E-21	7.67E-21	AAEL0000690-PA OS=Aedes aegypti GN=AAEL000690 PE=3 SV=1
CCG005486.1	1419	113	19	3.42183228	0.468081225	-2.869938245	Down	2.78E-22	1.11E-21	AAEL005004-PA OS=Aedes aegypti GN=AAEL005004 PE=4 SV=1
CCG018309.1	685	113	19	7.08843797	0.969645633	-2.869938245	Down	2.78E-22	1.11E-21	AAEL002221-PA OS=Aedes aegypti GN=AAEL002221 PE=4 SV=1
CCG001354.1	1125	139	19	5.30914943	0.590406452	-3.168700355	Down	9.00E-30	3.78E-29	AAEL010678-PA OS=Aedes aegypti GN=AAEL010678 PE=4 SV=1
CCG017896.1	668	167	19	10.7424337	0.994322244	-3.433463575	Down	4.09E-38	1.81E-37	-
CCG027756.1	1563	192	19	5.27843189	0.424956659	-3.634721783	Down	9.71E-46	4.45E-45	AAEL000863-PA OS=Aedes aegypti GN=AAEL000863 PE=4 SV=1
CCG013707.1	8613	215	19	1.07262196	0.07711683	-3.797952132	Down	7.28E-53	3.44E-52	AAEL000700-PA OS=Aedes aegypti GN=AAEL000700 PE=3 SV=1
CCG025368.1	1293	451	19	14.9878966	0.513694709	-4.866742906	Down	3.85E-129	2.14E-128	AAEL000624-PA OS=Aedes aegypti GN=AAEL006624 PE=4 SV=1
CCG025321.1	234	0	20	0.001	2.987886904	11.54490983	Up	7.45E-06	1.32E-05	AAEL007034-PA OS=Aedes aegypti GN=AAEL007034 PE=4 SV=1
CCG017177.1	387	0	20	0.001	1.806629291	10.81908479	Up	7.45E-06	1.32E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_22381 PE=4 SV=1
CCG016983.1	387	0	20	0.001	1.806629291	10.81908479	Up	7.45E-06	1.32E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011615 PE=4 SV=1
CCG013847.1	387	0	20	0.001	1.806629291	10.81908479	Up	7.45E-06	1.32E-05	AAEL001980-PA OS=Aedes aegypti GN=AAEL001980 PE=4 SV=1
CCG006777.1	660	0	20	0.001	1.05934172	10.04895233	Up	7.45E-06	1.32E-05	AAEL007525-PA OS=Aedes aegypti GN=AAEL007525 PE=3 SV=1
CCG003908.1	714	0	20	0.001	0.979223439	9.935494281	Up	7.45E-06	1.32E-05	-
CCG004194.1	728	0	20	0.001	0.960392219	9.907479905	Up	7.45E-06	1.32E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG018230.1	822	0	20	0.001	0.850566345	9.732279961	Up	7.45E-06	1.32E-05	AAEL001637-PA OS=Aedes aegypti GN=AAEL001637 PE=4 SV=1

CCG025315.1	905	0	20	0.001	0.772558603	9.593500563	Up	7.45E-06	1.32E-05	-
CCG009271.1	1686	0	20	0.001	0.41468893	8.695885724	Up	7.45E-06	1.32E-05	AAEL003201-PA (Fragment) OS=Aedes aegypti GN=AAEL003201 PE=4 SV=1
CCG013602.1	1854	0	20	0.001	0.377111939	8.558849016	Up	7.45E-06	1.32E-05	AAEL000603-PA (Fragment) OS=Aedes aegypti GN=AAEL000603 PE=4 SV=1
CCG015293.1	1866	2	20	0.04605545	0.374686782	3.024241299	Up	0.0005172	0.00081184	AAEL004778-PA OS=Aedes aegypti GN=AAEL004778 PE=3 SV=1
CCG006701.1	1521	2	20	0.05650195	0.459674908	3.024241299	Up	0.0005172	0.00081189	AAEL001237-PA (Fragment) OS=Aedes aegypti GN=AAEL001237 PE=4 SV=1
CCG003481.1	1157	2	20	0.07427785	0.604291733	3.024241299	Up	0.0005172	0.00081179	-
CCG015706.1	477	2	20	0.1801666	1.46575584	3.024241299	Up	0.0005172	0.00081194	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002557 PE=4 SV=1
CCG028471.2	714	41	20	2.46744975	0.979223439	-1.333310706	Down	0.0004667	0.00073485	AAEL005001-PA OS=Aedes aegypti GN=AAEL005001 PE=3 SV=1
CCG002390.1	1392	42	20	1.29650061	0.502274092	-1.368076124	Down	0.000303	0.00048206	RUNX2 OS=Aedes aegypti PE=2 SV=1
CCG012336.1	435	43	20	4.24758296	1.607277093	-1.402023456	Down	0.0001955	0.0003144	-
CCG011461.1	803	43	20	2.30099451	0.870691825	-1.402023456	Down	0.0001955	0.00031442	Retrotransposon protein, putative, unclassified OS=Oryza sativa subsp. japonica GN=LOC_Os12g44200 PE=4 SV=1
CCG007172.1	1291	44	20	1.46449909	0.541568966	-1.43519032	Down	0.0001253	0.00020355	-
CCG024393.1	651	46	20	3.03626389	1.073986998	-1.499320657	Down	5.05E-05	8.48E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013896 PE=4 SV=1
CCG002784.1	747	48	20	2.76110744	0.935964572	-1.560721202	Down	1.99E-05	3.43E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018758 PE=4 SV=1
CCG006912.1	1149	48	20	1.7950803	0.608499161	-1.560721202	Down	1.99E-05	3.43E-05	AAEL005748-PA OS=Aedes aegypti GN=AAEL005748 PE=4 SV=1

CCG018677.1	906	48	20	2.27654223	0.771705889	-1.560721202	Down	1.99E-05	3.43E-05	AAEL009792-PA OS=Aedes aegypti GN=AAEL009792 PE=4 SV=1
CCG008066.1	2435	49	20	0.8646887	0.287131637	-1.590468545	Down	1.24E-05	2.16E-05	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG019210.1	1068	58	20	2.33356237	0.654649378	-1.833739696	Down	1.40E-07	2.67E-07	AAEL005671-PA OS=Aedes aegypti GN=AAEL005671 PE=4 SV=1
CCG011807.1	2780	60	20	0.92740434	0.251498394	-1.882649297	Down	4.94E-08	9.61E-08	-
CCG022789.1	2502	66	20	1.1334942	0.27944266	-2.020152821	Down	1.98E-09	4.05E-09	Transporter OS=Aedes aegypti GN=AAEL005278 PE=3 SV=1
CCG005581.1	474	69	20	6.25508795	1.475032775	-2.084283158	Down	3.79E-10	7.91E-10	AAEL010187-PA (Fragment) OS=Aedes aegypti GN=AAEL010187 PE=4 SV=1
CCG011858.1	4401	69	20	0.67369045	0.158865152	-2.084283158	Down	3.79E-10	7.91E-10	AAEL008598-PA OS=Aedes aegypti GN=AAEL008598 PE=4 SV=1
CCG021619.1	2334	70	20	1.28872383	0.299556785	-2.105041718	Down	2.17E-10	4.58E-10	AAEL011203-PA OS=Aedes aegypti GN=AAEL011203 PE=4 SV=1
CCG000038.1	666	71	20	4.58085759	1.049798101	-2.125505821	Down	1.24E-10	2.64E-10	Gag polyprotein OS=Anopheles gambiae PE=4 SV=1
CCG007117.1	5409	72	20	0.5719765	0.129259666	-2.145683703	Down	7.06E-11	1.52E-10	AAEL006534-PA (Fragment) OS=Aedes aegypti GN=AAEL006534 PE=4 SV=1
CCG024766.1	2325	77	20	1.42308368	0.300716359	-2.242545242	Down	4.05E-12	9.36E-12	r-box/reducine rich repeat protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002568 PE=4 SV=1
CCG011017.1	1629	79	20	2.08386067	0.429199224	-2.279539449	Down	1.27E-12	3.09E-12	AAEL012524-PA OS=Aedes aegypti GN=AAEL012524 PE=4 SV=1
CCG025124.1	1341	80	20	2.56344427	0.521376238	-2.297686796	Down	7.08E-13	1.79E-12	AAEL011899-PA OS=Aedes aegypti GN=AAEL011899 PE=4 SV=1
CCG003304.1	717	80	20	4.79439159	0.97512627	-2.297686796	Down	7.08E-13	1.79E-12	AAEL009805-PA OS=Aedes aegypti GN=AAEL009805 PE=4 SV=1
CCG003912.1	495	84	20	7.29183375	1.412455627	-2.368076124	Down	6.70E-14	2.20E-13	GF13847 OS=Drosophila ananassae GN=GF13847 PE=4 SV=1
CCG011147.1	901	89	20	4.24451319	0.775988386	-2.451492132	Down	3.35E-15	1.21E-14	-

CCG009778.1	927	90	20	4.17181889	0.754223879	-2.467611798	Down	1.83E-15	6.63E-15	AAEL005779-PA OS=Aedes aegypti GN=AAEL005779 PE=4 SV=1
CCG024903.1	1294	96	20	3.18786285	0.540313397	-2.560721202	Down	4.65E-17	1.76E-16	AAEL000372-PA OS=Aedes aegypti GN=AAEL000372 PE=4 SV=1
CCG002673.1	720	97	20	5.78897814	0.971063244	-2.575671543	Down	2.51E-17	9.52E-17	AAEL011202-PA OS=Aedes aegypti GN=AAEL011202 PE=4 SV=1
CCG011196.1	1135	98	20	3.71016211	0.616004877	-2.590468545	Down	1.35E-17	5.14E-17	Ubiquitin carboxyl-terminal hydrolase OS=Culex quinquefasciatus GN=CpipJ_CPIJ013438 PE=3 SV=1
CCG019751.1	1473	113	20	3.29638833	0.474654131	-2.795937664	Down	1.04E-21	4.10E-21	AAEL009314-PA OS=Aedes aegypti GN=AAEL009314 PE=4 SV=1
CCG014807.1	912	134	20	6.31353557	0.766628877	-3.041847892	Down	1.12E-27	4.65E-27	AAEL005989-PA OS=Aedes aegypti GN=AAEL005989 PE=4 SV=1
CCG005535.1	1830	159	20	3.73343596	0.38205767	-3.288641657	Down	5.21E-35	2.26E-34	AAEL005345-PA OS=Aedes aegypti GN=AAEL005345 PE=4 SV=1
CCG001433.1	1170	161	20	5.91292929	0.597577381	-3.306675579	Down	1.32E-35	5.75E-35	AAEL006027-PA (Fragment) OS=Aedes aegypti GN=AAEL006027 PE=3 SV=1
CCG010948.1	1494	183	20	5.26336107	0.467982286	-3.49145854	Down	3.16E-42	1.43E-41	AAEL014332-PA OS=Aedes aegypti GN=AAEL014332 PE=4 SV=1
CCG009348.1	682	190	20	11.9710404	1.025169407	-3.54561431	Down	2.34E-44	1.06E-43	AAEL012693-PA OS=Aedes aegypti GN=AAEL012693 PE=4 SV=1
CCG018040.1	696	191	20	11.7919818	1.004548183	-3.553187529	Down	1.16E-44	5.28E-44	AAEL010397-PA OS=Aedes aegypti GN=AAEL010397 PE=4 SV=1
CCG024052.1	3708	261	20	3.0245687	0.18855597	-4.003664698	Down	1.84E-66	9.00E-66	AAEL015231-PA OS=Aedes aegypti GN=AAEL015231 PE=4 SV=1
CCG027137.1	294	262	20	38.2927567	2.378114066	-4.009181703	Down	8.86E-67	4.35E-66	-
CCG019654.1	783	332	20	18.2196065	0.892931718	-4.350798133	Down	3.56E-89	1.85E-88	AAEL002738-PA (Fragment) OS=Aedes aegypti GN=AAEL002738 PE=4 SV=1
CCG015672.1	543	446	20	35.2937415	1.287597671	-4.776658601	Down	2.22E-126	1.23E-125	AAEL011068-PA OS=Aedes aegypti GN=AAEL011068 PE=4 SV=1
CCG006318.2	2014	530	20	11.3078249	0.347152699	-5.025607251	Down	3.67E-154	2.10E-153	AAEL006334-PA OS=Aedes aegypti GN=AAEL006334 PE=4 SV=1

CCG016571.1	597	33210	20	2390.32644	1.17113155	-10.9950888	Down	0	0	AAEL010180-PA OS=Aedes aegypti GN=AAEL010180 PE=4 SV=1
CCG026761.1	633	0	21	0.001	1.159753258	10.17960218	Up	4.11E-06	7.35E-06	AAEL010402-PA OS=Aedes aegypti GN=AAEL010402 PE=3 SV=1
CCG000073.1	824	0	21	0.001	0.890926957	9.799163346	Up	4.11E-06	7.34E-06	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010212 PE=4 SV=1
CCG013494.1	836	0	21	0.001	0.878138531	9.778304741	Up	4.11E-06	7.35E-06	AAEL005156-PA OS=Aedes aegypti GN=AAEL005156 PE=4 SV=1
CCG025336.1	975	0	21	0.001	0.7529475	9.556405464	Up	4.11E-06	7.35E-06	AAEL001237-PA (Fragment) OS=Aedes aegypti GN=AAEL001237 PE=4 SV=1
CCG008479.1	1017	0	21	0.001	0.721852323	9.495559909	Up	4.11E-06	7.34E-06	AAEL007070-PA OS=Aedes aegypti GN=AAEL007070 PE=3 SV=1
CCG006699.1	4227	0	21	0.001	0.173674902	7.440245476	Up	4.11E-06	7.35E-06	AAEL002654-PA OS=Aedes aegypti GN=AAEL002654 PE=4 SV=1
CCG011750.1	891	1	21	0.04822641	0.823932449	4.094630627	Up	0.0006316	0.00098158	-
CCG000734.2	465	1	21	0.09240803	1.578760886	4.094630627	Up	0.0006316	0.00098164	-
CCG016336.1	777	2	21	0.11060421	0.944818291	3.094630627	Up	0.0003472	0.0005506	AAEL010959-PA OS=Aedes aegypti GN=AAEL010959 PE=3 SV=1
CCG006544.1	363	2	21	0.23674785	2.022379648	3.094630627	Up	0.0003472	0.00055056	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001635 PE=4 SV=1
CCG018932.1	1002	2	21	0.08576793	0.732658495	3.094630627	Up	0.0003472	0.00055049	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009045 PE=4 SV=1
CCG013642.1	354	2	21	0.24276686	2.07379608	3.094630627	Up	0.0003472	0.00055046	AAEL009961-PA OS=Aedes aegypti GN=AAEL009961 PE=4 SV=1
CCG025109.1	432	2	21	0.19893396	1.699360676	3.094630627	Up	0.0003472	0.00055053	-
CCG006779.1	1479	42	21	1.22023587	0.496364985	-1.297686796	Down	0.0005233	0.00082113	AAEL007531-PA OS=Aedes aegypti GN=AAEL007531 PE=4 SV=1
CCG020046.1	521	46	21	3.79387292	1.409066818	-1.428931329	Down	9.24E-05	0.0001523	-

CCG021569.1	430	49	21	4.89655115	1.70726468	-1.520079218	Down	2.36E-05	4.06E-05	AAEL004044-PA (Fragment) OS=Aedes aegypti GN=AAEL004044 PE=4 SV=1
CCG013366.1	2760	52	21	0.80957471	0.265986888	-1.605809092	Down	5.76E-06	1.02E-05	AAEL014568-PA OS=Aedes aegypti GN=AAEL014568 PE=3 SV=1
CCG019563.1	1221	73	21	2.56903409	0.601248003	-2.095193932	Down	1.01E-10	2.16E-10	-
CCG022211.1	909	80	21	3.78171482	0.807616955	-2.227297468	Down	1.92E-12	4.59E-12	Angiotensin 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017837 PE=4 SV=1
CCG013086.1	588	88	21	6.43084464	1.248509885	-2.364800992	Down	1.79E-14	6.28E-14	AAEL008440-PA OS=Aedes aegypti GN=AAEL008440 PE=4 SV=1
CCG008185.1	1617	96	21	2.55107886	0.454003594	-2.490331874	Down	1.46E-16	5.48E-16	AAEL011709-PA (Fragment) OS=Aedes aegypti GN=AAEL011709 PE=4 SV=1
CCG018350.1	1659	113	21	2.92681134	0.442509833	-2.725548336	Down	3.70E-21	1.46E-20	AAEL008628-PA OS=Aedes aegypti GN=AAEL008628 PE=3 SV=1
CCG016072.1	2232	124	21	2.38720748	0.328908518	-2.859565684	Down	3.19E-24	1.29E-23	Carboxypeptidase B OS=Aedes aegypti GN=CPB-II PE=2 SV=1
CCG013148.1	948	173	21	7.8415233	0.774392207	-3.339997601	Down	1.72E-38	7.64E-38	AAEL000715-PA OS=Aedes aegypti GN=AAEL000715 PE=4 SV=1
CCG002839.1	1299	179	21	5.92115666	0.565145352	-3.389185151	Down	2.74E-40	1.22E-39	AAEL003958-PA OS=Aedes aegypti GN=AAEL003958 PE=3 SV=1
CCG012215.1	993	227	21	9.82288999	0.739298905	-3.731917861	Down	5.80E-55	2.75E-54	AAEL011454-PA OS=Aedes aegypti GN=AAEL011454 PE=4 SV=1
CCG012285.1	954	239	21	10.7649545	0.769521816	-3.806236181	Down	1.07E-58	5.15E-58	AAEL014757-PA OS=Aedes aegypti GN=AAEL014757 PE=4 SV=1
CCG015492.1	666	316	21	20.3880422	1.102288006	-4.209150122	Down	4.55E-83	2.32E-82	AAEL005777-PA OS=Aedes aegypti GN=AAEL005777 PE=4 SV=1
CCG018048.1	1248	2201	21	75.7823605	0.588240234	-7.009312803	Down	0	0	AAEL005969-PA OS=Aedes aegypti GN=AAEL005969 PE=4 SV=1
CCG000318.1	297	0	22	0.001	2.589501983	11.33845895	Up	2.26E-06	4.09E-06	-
CCG005904.1	371	0	22	0.001	2.072997544	11.01750269	Up	2.26E-06	4.09E-06	AAEL010586-PA OS=Aedes aegypti GN=AAEL010586 PE=4 SV=1

CCG006761.1	456	0	22	0.001	1.686583528	10.71988805	Up	2.26E-06	4.09E-06	AGAP003907-PA OS=Anopheles gambiae GN=AGAP003907 PE=4 SV=2
CCG003228.1	573	0	22	0.001	1.342202599	10.39038674	Up	2.26E-06	4.09E-06	modifer of mdg4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001640 PE=4 SV=1
CCG005663.1	579	0	22	0.001	1.328293763	10.37535853	Up	2.26E-06	4.09E-06	AAEL013351-PA OS=Aedes aegypti GN=AAEL013351 PE=3 SV=1
CCG018313.1	1032	0	22	0.001	0.745234582	9.541550813	Up	2.26E-06	4.09E-06	AAEL003984-PA OS=Aedes aegypti GN=AAEL003984 PE=4 SV=1
CCG002259.1	425	1	22	0.10110526	1.809604915	4.161744822	Up	0.0006126	0.00095276	Uncharacterized protein OS=Metarhizium anisopliae (strain ARSEF 23 / ATCC MYA-3075) GN=MAA_00416 PE=4 SV=1
CCG018950.1	435	1	22	0.098781	1.768004802	4.161744822	Up	0.0006126	0.0009527	AAEL006494-PA (Fragment) OS=Aedes aegypti GN=GPRgr45 PE=3 SV=2
CCG000678.1	456	1	22	0.09423187	1.686583528	4.161744822	Up	0.0006126	0.00095264	AAEL013835-PA OS=Aedes aegypti GN=AAEL013835 PE=4 SV=1
CCG003689.1	933	1	22	0.04605545	0.824310921	4.161744822	Up	0.0006126	0.00095258	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008885 PE=4 SV=1
CCG017166.1	1862	2	22	0.04615439	0.413040864	3.161744822	Up	0.0002449	0.00039213	RNase H and integrase-like protein (Fragment) OS=Bombyx mori GN=P-1 PE=4 SV=1
CCG022134.1	438	2	22	0.19620883	1.75589518	3.161744822	Up	0.0002449	0.00039216	AGAP000591-PA OS=Anopheles gambiae GN=AgaP_AGAP000591 PE=3 SV=4
CCG013782.1	1263	44	22	1.49696621	0.60893277	-1.297686796	Down	0.000384	0.00060744	AAEL007784-PA OS=Aedes aegypti GN=AAEL007784 PE=4 SV=1
CCG006444.1	2856	44	22	0.66199871	0.269286446	-1.297686796	Down	0.000384	0.0006074	AAEL009774-PA OS=Aedes aegypti GN=AAEL009774 PE=3 SV=1
CCG023002.1	1941	46	22	1.01834508	0.396229824	-1.361817134	Down	0.0001638	0.0002643	AAEL005681-PA OS=Aedes aegypti GN=GPRHIS PE=4 SV=2

CCG012201.1	1554	47	22	1.29959944	0.494904819	-1.392844029	Down	0.000106	0.0001728	Putative protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012062 PE=4 SV=1
CCG024683.1	1479	48	22	1.39455528	0.520001412	-1.423217678	Down	6.82E-05	0.00011319	AAEL007260-PA OS=Aedes aegypti GN=AAEL007260 PE=4 SV=1
CCG022496.1	474	48	22	4.35136553	1.622536053	-1.423217678	Down	6.82E-05	0.0001132	AGAP003669-PA OS=Anopheles gambiae GN=AgaP_AGAP003669 PE=3 SV=4
CCG014575.1	1437	49	22	1.46521712	0.535199784	-1.452965022	Down	4.36E-05	7.35E-05	AAEL015017-PA OS=Aedes aegypti GN=AAEL015017 PE=4 SV=1
CCG017230.3	1620	53	22	1.40579996	0.47474203	-1.566175632	Down	6.92E-06	1.23E-05	AGAP006349-PA OS=Anopheles gambiae GN=AGAP006349 PE=4 SV=4
CCG028639.1	652	56	22	3.69065205	1.179573756	-1.6456101	Down	1.65E-06	3.01E-06	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG020777.1	614	57	22	3.98904702	1.252576692	-1.671145192	Down	1.01E-06	1.87E-06	-
CCG013966.1	1986	59	22	1.27654297	0.387251807	-1.720898227	Down	3.78E-07	7.08E-07	AAEL006452-PA OS=Aedes aegypti GN=AAEL006452 PE=3 SV=1
CCG011706.1	4512	60	22	0.57140605	0.170452591	-1.745145773	Down	2.29E-07	4.33E-07	Ion channel nompC OS=Culex quinquefasciatus GN=CpipJ_CPIJ000429 PE=4 SV=1
CCG009023.1	2301	61	22	1.13913681	0.334238196	-1.768992515	Down	1.39E-07	2.64E-07	AAEL014203-PA OS=Aedes aegypti GN=AAEL014203 PE=4 SV=1
CCG010081.1	582	63	22	4.65136302	1.321446888	-1.815535101	Down	5.00E-08	9.71E-08	AAEL004494-PA (Fragment) OS=Aedes aegypti GN=AAEL004494 PE=4 SV=1
CCG007626.1	1534	66	22	1.84876303	0.501357294	-1.882649297	Down	1.05E-08	2.10E-08	AAEL002133-PA OS=Aedes aegypti GN=AAEL002133 PE=4 SV=1
CCG006020.1	2555	68	22	1.1436172	0.301010602	-1.925718019	Down	3.66E-09	7.41E-09	RNase H and integrase-like protein (Fragment) OS=Bombyx mori GN=P-12 PE=4 SV=1
CCG017496.1	1059	69	22	2.79972775	0.726234267	-1.946779634	Down	2.15E-09	4.39E-09	AGAP000277-PA OS=Anopheles gambiae GN=AgaP_AGAP000277 PE=4 SV=2

CCG014988.1	597	77	22	5.54216008	1.288244705	-2.105041718	Down	2.71E-11	5.90E-11	AAEL001355-PA OS=Aedes aegypti GN=AAEL001355 PE=4 SV=1
CCG018863.1	1191	77	22	2.77806009	0.645744827	-2.105041718	Down	2.71E-11	5.90E-11	AAEL011874-PA OS=Aedes aegypti GN=AAEL011874 PE=4 SV=1
CCG012189.1	2367	85	22	1.54306187	0.3249185	-2.247646114	Down	2.87E-13	7.98E-13	AAEL014297-PA OS=Aedes aegypti GN=AAEL014297 PE=4 SV=1
CCG025058.1	1566	102	22	2.79879497	0.491112445	-2.51068052	Down	1.16E-17	4.41E-17	AAEL009546-PA OS=Aedes aegypti GN=AAEL009546 PE=4 SV=1
CCG025084.1	1516	140	22	3.9681813	0.507310085	-2.967538194	Down	3.60E-28	1.50E-27	AAEL008168-PA OS=Aedes aegypti GN=AAEL008168 PE=4 SV=1
CCG007371.2	2763	148	22	2.30167236	0.278350376	-3.047708543	Down	1.81E-30	7.63E-30	AAEL007967-PA OS=Aedes aegypti GN=AAEL007967 PE=4 SV=1
CCG001038.1	1056	149	22	6.06296445	0.728297433	-3.057423698	Down	9.29E-31	3.93E-30	Galactose-1-phosphate uridylyltransferase OS=Aedes aegypti GN=AAEL001607 PE=3 SV=1
CCG028022.1	1775	151	22	3.65545348	0.433285684	-3.076659917	Down	2.45E-31	1.04E-30	AAEL006248-PA OS=Aedes aegypti GN=AAEL006248 PE=4 SV=1
CCG015646.1	2001	168	22	3.60765388	0.38434887	-3.2305726	Down	2.58E-36	1.13E-35	AAEL009258-PA OS=Aedes aegypti GN=AAEL009258 PE=4 SV=1
CCG016997.1	1662	310	22	8.01481211	0.462744939	-4.114379583	Down	3.17E-80	1.61E-79	AGAP007672-PA OS=Anopheles gambiae GN=AgaP_AGAP007672 PE=3 SV=5
CCG011876.1	1371	344	22	10.781611	0.560964325	-4.264519932	Down	4.17E-91	2.17E-90	AAEL003596-PA OS=Aedes aegypti GN=AAEL003596 PE=4 SV=1
CCG001335.1	1137	379	22	14.3232449	0.676413447	-4.404309216	Down	2.13E-102	1.13E-101	AAEL005530-PA OS=Aedes aegypti GN=AAEL005530 PE=4 SV=1
CCG004861.1	1185	699	22	25.3467042	0.649014421	-5.287403823	Down	4.18E-208	2.50E-207	AAEL001437-PA OS=Aedes aegypti GN=AAEL001437 PE=4 SV=1
CCG024829.1	2865	1125	22	16.8729324	0.26844052	-5.973964464	Down	0	0	AAEL004897-PA OS=Aedes aegypti GN=AAEL004897 PE=4 SV=1
CCG002498.1	576	5062	22	377.626383	1.33521196	-8.14374697	Down	0	0	AAEL001677-PA OS=Aedes aegypti GN=AAEL001677 PE=4 SV=1
CCG020332.1	273	0	23	0.001	2.945202805	11.52415127	Up	1.25E-06	2.29E-06	-

CCG004746.1	515	0	23	0.001	1.561243429	10.60847978	Up	1.25E-06	2.29E-06	tetratricopeptide repeat protein, tpr OS=Culex quinquefasciatus GN=CpipJ_CPIJ016841 PE=4 SV=1
CCG027213.1	549	0	23	0.001	1.4645544	10.51624607	Up	1.25E-06	2.29E-06	AAEL010722-PA OS=Aedes aegypti GN=AAEL010722 PE=4 SV=1
CCG018503.1	1089	0	23	0.001	0.738329078	9.528120168	Up	1.25E-06	2.29E-06	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG015525.1	1938	0	23	0.001	0.41488151	8.696555551	Up	1.25E-06	2.29E-06	AAEL004028-PA OS=Aedes aegypti GN=AAEL004028 PE=3 SV=1
CCG006816.1	921	1	23	0.04665552	0.873007998	4.22587516	Up	0.0006016	0.00093599	AAEL004556-PA OS=Aedes aegypti GN=AAEL004556 PE=4 SV=1
CCG006460.1	528	1	23	0.08138207	1.522803723	4.22587516	Up	0.0006016	0.00093605	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003878 PE=4 SV=1
CCG025567.1	648	2	23	0.13262264	1.240803034	3.22587516	Up	0.0001836	0.00029566	-
CCG016096.1	204	2	23	0.42127191	3.941374342	3.22587516	Up	0.0001836	0.00029562	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014565 PE=4 SV=1
CCG006935.1	1371	2	23	0.06268378	0.586462703	3.22587516	Up	0.0001836	0.00029558	AAEL010721-PA OS=Aedes aegypti GN=AAEL010721 PE=4 SV=1
CCG019990.1	849	2	23	0.10122435	0.947044011	3.22587516	Up	0.0001836	0.0002956	AAEL013652-PA OS=Aedes aegypti GN=AAEL013652 PE=3 SV=1
CCG028594.1	915	2	23	0.09392292	0.87873264	3.22587516	Up	0.0001836	0.00029564	AAEL006410-PA OS=Aedes aegypti GN=AAEL006410 PE=4 SV=1
CCG023081.1	621	3	23	0.20758326	1.294750992	2.640912659	Up	0.0004358	0.00068724	AAEL002330-PA OS=Aedes aegypti GN=AAEL002330 PE=4 SV=1
CCG014696.1	2685	3	23	0.04801088	0.299456375	2.640912659	Up	0.0004358	0.00068742	Polyprotein OS=Antheraea mylitta PE=4 SV=1
CCG005736.1	828	3	23	0.15568744	0.971063244	2.640912659	Up	0.0004358	0.00068737	AAEL005100-PA OS=Aedes aegypti GN=AAEL005100 PE=4 SV=1
CCG003624.1	2220	3	23	0.05806721	0.362180345	2.640912659	Up	0.0004358	0.00068733	AAEL015386-PA OS=Aedes aegypti GN=AAEL015386 PE=4 SV=1
CCG003289.1	1632	3	23	0.07898848	0.492671793	2.640912659	Up	0.0004358	0.00068729	AAEL004091-PA (Fragment) OS=Aedes aegypti GN=AAEL004091 PE=3 SV=1

CCG015263.1	867	47	23	2.32938584	0.927382198	-1.328713692	Down	0.0001849	0.00029772	AAEL004964-PA OS=Aedes aegypti GN=AAEL004964 PE=4 SV=1
CCG004000.1	1116	48	23	1.84816063	0.720466278	-1.359087341	Down	0.0001206	0.00019605	AAEL002726-PA OS=Aedes aegypti GN=AAEL002726 PE=4 SV=1
CCG013388.1	1992	52	23	1.1216999	0.403634722	-1.474564558	Down	2.06E-05	3.54E-05	AAEL012755-PA OS=Aedes aegypti GN=AAEL012755 PE=4 SV=1
CCG010333.1	2268	55	23	1.04203501	0.354515152	-1.555484554	Down	5.17E-06	9.18E-06	AAEL013476-PA OS=Aedes aegypti GN=AAEL013476 PE=3 SV=1
CCG015320.1	663	56	23	3.62941951	1.212730567	-1.581479762	Down	3.23E-06	5.79E-06	-
CCG022508.1	1227	56	23	1.96112888	0.655289622	-1.581479762	Down	3.23E-06	5.79E-06	-
CCG016868.1	633	62	23	4.20872598	1.270205949	-1.728321151	Down	1.75E-07	3.32E-07	AAEL015308-PA (Fragment) OS=Aedes aegypti GN=SCRAL1 PE=4 SV=2
CCG004969.1	1503	70	23	2.00125178	0.534956997	-1.903407857	Down	2.88E-09	5.86E-09	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011425 PE=4 SV=1
CCG027051.1	1323	76	23	2.46840501	0.607740261	-2.022052354	Down	1.15E-10	2.45E-10	AAEL007016-PA OS=Aedes aegypti GN=AAEL007016 PE=3 SV=1
CCG009236.1	1617	81	23	2.15247279	0.497242032	-2.113974843	Down	7.25E-12	1.63E-11	Beta-1,3-glucan-binding protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013556 PE=4 SV=1
CCG018282.1	966	93	23	4.1368378	0.832339923	-2.313283651	Down	7.47E-15	2.68E-14	AAEL014627-PA OS=Aedes aegypti GN=AAEL014627 PE=3 SV=1
CCG007530.1	1384	95	23	2.94951213	0.580954022	-2.343980448	Down	2.30E-15	8.34E-15	Uncharacterized protein OS=Clonorchis sinensis GN=CLF_109243 PE=4 SV=1
CCG006543.1	456	102	23	9.61165116	1.763246416	-2.446550182	Down	3.53E-17	1.34E-16	AGAP006213-PA OS=Anopheles gambiae GN=AGAP006213 PE=4 SV=4
CCG024667.1	1011	114	23	4.84525197	0.795292152	-2.607014854	Down	2.27E-20	8.87E-20	-
CCG000760.1	3174	115	23	1.55687444	0.253320846	-2.619614891	Down	1.22E-20	4.77E-20	AAEL009473-PA OS=Aedes aegypti GN=AAEL009473 PE=4 SV=1
CCG004006.1	1338	136	23	4.36762624	0.60092703	-2.861587681	Down	1.92E-26	7.89E-26	AAEL010315-PA OS=Aedes aegypti GN=AAEL010315 PE=4 SV=1

CCG027037.1	1483	187	23	5.41830099	0.542171521	-3.3210193	Down	2.85E-41	1.28E-40	AAEL006334-PA OS=Aedes aegypti GN=AAEL006334 PE=4 SV=1
CCG014134.1	386	409	23	45.5301074	2.083006129	-4.450081873	Down	4.02E-111	2.17E-110	AAEL001504-PA OS=Aedes aegypti GN=AAEL001504 PE=4 SV=1
CCG028320.1	1221	488	23	17.1738169	0.658509718	-4.704862178	Down	6.49E-137	3.64E-136	AAEL000044-PA OS=Aedes aegypti GN=CYP325Q1 PE=3 SV=1
CCG005493.1	2817	1909	23	29.1193551	0.28542434	-6.672726228	Down	0	0	AAEL001953-PA OS=Aedes aegypti GN=AAEL001953 PE=4 SV=1
CCG000671.1	433	0	24	0.001	1.937641207	10.92008574	Up	6.88E-07	1.27E-06	Nucleosome-remodeling factor subunit NURF301 OS=Culex quinquefasciatus GN=CpipJ_CPIJ011284 PE=4 SV=1
CCG019076.1	522	0	24	0.001	1.607277093	10.65040295	Up	6.88E-07	1.27E-06	AGAP004620-PC OS=Anopheles gambiae GN=AgaP_AGAP004620 PE=3 SV=1
CCG000644.1	1263	0	24	0.001	0.664290295	9.375670027	Up	6.88E-07	1.27E-06	AAEL012042-PA (Fragment) OS=Aedes aegypti GN=AAEL012042 PE=3 SV=1
CCG002292.1	2073	0	24	0.001	0.404726793	8.66080455	Up	6.88E-07	1.27E-06	AAEL015078-PA OS=Aedes aegypti GN=AAEL015078 PE=4 SV=1
CCG022023.1	634	1	24	0.06777561	1.323341707	4.287275705	Up	0.0005954	0.00092668	-
CCG006414.1	533	1	24	0.08061864	1.574106271	4.287275705	Up	0.0005954	0.00092685	-
CCG020550.1	513	1	24	0.08376167	1.635474937	4.287275705	Up	0.0005954	0.0009268	-
CCG007667.1	3242	1	24	0.01325408	0.258790451	4.287275705	Up	0.0005954	0.00092674	Elongation factor 1-alpha OS=Aedes aegypti PE=2 SV=1
CCG012172.1	410	2	24	0.20960846	2.046338153	3.287275705	Up	0.000147	0.00023784	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001647 PE=4 SV=1
CCG015815.1	825	2	24	0.10416905	1.016968052	3.287275705	Up	0.000147	0.00023779	AAEL012564-PA OS=Aedes aegypti GN=AAEL012564 PE=3 SV=1
CCG002416.1	1687	2	24	0.05094219	0.497331738	3.287275705	Up	0.000147	0.00023782	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG021794.1	297	2	24	0.28935848	2.824911254	3.287275705	Up	0.000147	0.00023781	AAEL005936-PA OS=Aedes aegypti GN=AAEL005936 PE=4 SV=1
CCG019917.1	687	2	24	0.12509384	1.221249844	3.287275705	Up	0.000147	0.00023778	-

CCG017053.1	1176	3	24	0.10961667	0.71343422	2.702313204	Up	0.000288	0.00045888	Uncharacterized protein OS=Anopheles darlingi GN=AND_17550 PE=4 SV=1
CCG003558.1	1471	3	24	0.08763372	0.570359376	2.702313204	Up	0.000288	0.00045891	AGAP001115-PA OS=Anopheles gambiae GN=GPRGR42 PE=3 SV=2
CCG018413.1	2788	3	24	0.04623716	0.300932081	2.702313204	Up	0.000288	0.00045885	-
CCG021540.1	930	46	24	2.12538472	0.902149078	-1.236286252	Down	0.0004719	0.00074269	methionine-s-methyltransferase dehydrogenase, mitochondrial OS=Culex quinquefasciatus GN=CpipJ_CPIJ009984 PE=3 SV=1
CCG007324.1	414	48	24	4.98199822	2.026566769	-1.297686796	Down	0.0002072	0.00033281	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009250 PE=4 SV=1
CCG007235.1	633	49	24	3.32625118	1.325432295	-1.32743414	Down	0.000136	0.00022051	AAEL000571-PA OS=Aedes aegypti GN=AAEL000571 PE=4 SV=1
CCG006071.1	1311	51	24	1.67159151	0.639968453	-1.385149637	Down	5.76E-05	9.60E-05	AAEL007057-PA (Fragment) OS=Aedes aegypti GN=AAEL007057 PE=4 SV=1
CCG013913.1	1668	52	24	1.33958405	0.502996788	-1.413164014	Down	3.72E-05	6.28E-05	AAEL002775-PA OS=Aedes aegypti GN=AAEL002775 PE=4 SV=1
CCG013354.1	1008	53	24	2.25932136	0.832339923	-1.44064475	Down	2.39E-05	4.10E-05	-
CCG021629.1	1295	53	24	1.7586069	0.6478754	-1.44064475	Down	2.39E-05	4.10E-05	-
CCG002041.1	1519	64	24	1.81044306	0.55233617	-1.712724295	Down	1.33E-07	2.53E-07	-
CCG011895.1	1917	65	24	1.45698109	0.437662307	-1.735092108	Down	8.09E-08	1.56E-07	AAEL000558-PA OS=Aedes aegypti GN=CYP325E3 PE=3 SV=1
CCG003145.1	491	76	24	6.65111982	1.708754873	-1.960651809	Down	2.70E-10	5.69E-10	AAEL008001-PA OS=Aedes aegypti GN=AAEL008001 PE=4 SV=1
CCG006403.1	618	77	24	5.35383425	1.357602981	-1.979510836	Down	1.58E-10	3.36E-10	AAEL008489-PA OS=Aedes aegypti GN=AAEL008489 PE=2 SV=1
CCG011731.1	468	86	24	7.89614781	1.792732142	-2.13898905	Down	1.11E-12	2.72E-12	-
CCG015740.1	2065	94	24	1.95600729	0.406294742	-2.267313147	Down	1.15E-14	4.10E-14	-
CCG011252.1	375	98	24	11.229424	2.237329713	-2.32743414	Down	1.11E-15	4.03E-15	AAEL006281-PA (Fragment) OS=Aedes aegypti GN=AAEL006281 PE=3 SV=1

CCG003272.1	3579	109	24	1.30866194	0.234422644	-2.48090862	Down	1.55E-18	5.95E-18	Uncharacterized protein OS=Anopheles darlingi GN=AND_23101 PE=4 SV=1
CCG012508.1	696	118	24	7.28509868	1.20545782	-2.595367345	Down	6.17E-21	2.43E-20	AAEL010918-PA OS=Aedes aegypti GN=AAEL010918 PE=4 SV=1
CCG008871.1	889	146	24	7.0568968	0.943755503	-2.902548854	Down	1.11E-28	4.63E-28	AAEL009225-PA OS=Aedes aegypti GN=AAEL009225 PE=4 SV=1
CCG021029.1	1410	171	24	5.21122313	0.595034498	-3.13057681	Down	7.10E-36	3.09E-35	Lipase OS=Aedes aegypti GN=AAEL014919 PE=3 SV=1
CCG028273.1	736	186	24	10.8591992	1.139943808	-3.251883107	Down	2.76E-40	1.23E-39	AAEL001396-PA OS=Aedes aegypti GN=AAEL001396 PE=4 SV=1
CCG004186.1	1629	212	24	5.59213244	0.515039068	-3.44064475	Down	4.58E-48	2.12E-47	AAEL012524-PA OS=Aedes aegypti GN=AAEL012524 PE=4 SV=1
CCG025836.1	5172	265	24	2.20165887	0.162219382	-3.762572845	Down	2.53E-64	1.23E-63	AAEL002204-PA OS=Aedes aegypti GN=AAEL002204 PE=4 SV=1
CCG015757.1	2372	279	24	5.05419728	0.353709377	-3.836845607	Down	1.09E-68	5.40E-68	-
CCG012208.1	453	279	24	26.4648034	1.852094134	-3.836845607	Down	1.09E-68	5.40E-68	Superoxide dismutase [Cu-Zn] OS=Aedes aegypti GN=CUSOD4 PE=3 SV=2
CCG026579.1	981	307	24	13.4472054	0.855248361	-3.974819141	Down	1.74E-77	8.79E-77	AAEL000432-PA OS=Aedes aegypti GN=AAEL000432 PE=3 SV=1
CCG001578.1	1001	434	24	18.6302346	0.838160482	-4.474275528	Down	3.15E-118	1.72E-117	AAEL004257-PA (Fragment) OS=Aedes aegypti GN=AAEL004257 PE=4 SV=1
CCG015804.1	2952	584	24	8.50078761	0.284213632	-4.902548854	Down	1.93E-167	1.12E-166	AAEL007483-PA OS=Aedes aegypti GN=AAEL007483 PE=4 SV=1
CCG008001.1	951	638	24	28.8272247	0.882227805	-5.030136909	Down	2.44E-185	1.44E-184	AAEL004353-PA OS=Aedes aegypti GN=AAEL004353 PE=4 SV=1
CCG004155.2	372	654	24	75.5435657	2.255372695	-5.065871121	Down	1.17E-190	6.95E-190	AAEL004829-PA OS=Aedes aegypti GN=AAEL004829 PE=4 SV=1
CCG018722.1	1209	977	24	34.7240949	0.693960829	-5.644939047	Down	5.35E-299	3.37E-298	AAEL001648-PA (Fragment) OS=Aedes aegypti GN=AAEL001648 PE=3 SV=1
CCG028030.1	531	0	25	0.001	1.645869905	10.68463459	Up	3.80E-07	7.11E-07	-

CCG017274.1	647	0	25	0.001	1.350783492	10.39958074	Up	3.80E-07	7.11E-07	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG011964.1	702	0	25	0.001	1.244952877	10.28187542	Up	3.80E-07	7.11E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000496 PE=4 SV=1
CCG024471.1	1170	0	25	0.001	0.746971726	9.544909825	Up	3.80E-07	7.11E-07	AAEL006371-PA OS=Aedes aegypti GN=AAEL006371 PE=3 SV=1
CCG006963.1	1908	0	25	0.001	0.4580487	8.839357184	Up	3.80E-07	7.11E-07	AAEL006375-PA OS=Aedes aegypti GN=AAEL006375 PE=4 SV=1
CCG009344.1	1554	1	25	0.02765105	0.56239184	4.346169394	Up	0.0005918	0.00092136	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001210 PE=4 SV=1
CCG028585.1	609	1	25	0.07055786	1.435068833	4.346169394	Up	0.0005918	0.00092153	-
CCG014490.1	497	1	25	0.08645822	1.758464626	4.346169394	Up	0.0005918	0.00092148	-
CCG015051.1	840	1	25	0.05115445	1.040424904	4.346169394	Up	0.0005918	0.00092142	AAEL013780-PA OS=Aedes aegypti GN=AAEL013780 PE=4 SV=1
CCG018656.1	836	2	25	0.10279841	1.045403014	3.346169394	Up	0.0001252	0.0002034	AAEL005607-PA OS=Aedes aegypti GN=AAEL005607 PE=3 SV=1
CCG004492.1	838	2	25	0.10255307	1.042908018	3.346169394	Up	0.0001252	0.00020337	-
CCG018437.1	1701	2	25	0.05052291	0.513790076	3.346169394	Up	0.0001252	0.00020338	AAEL008252-PA OS=Aedes aegypti GN=AAEL008252 PE=3 SV=1
CCG022719.1	723	3	25	0.17829765	1.20879242	2.761206893	Up	0.0001966	0.00031621	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008144 PE=4 SV=1
CCG024386.1	2223	3	25	0.05798885	0.393143014	2.761206893	Up	0.0001966	0.00031623	AAEL002752-PA OS=Aedes aegypti GN=AAEL002752 PE=4 SV=1
CCG003050.1	820	3	25	0.15720635	1.065801121	2.761206893	Up	0.0001966	0.00031619	AGAP003439-PL OS=Anopheles gambiae GN=mod(mdg4) PE=4 SV=1
CCG022424.1	854	4	25	0.20126339	1.023368758	2.346169394	Up	0.0005384	0.00084409	-
CCG003768.1	528	4	25	0.32552829	1.655221438	2.346169394	Up	0.0005384	0.00084403	AAEL009244-PA OS=Aedes aegypti GN=AAEL009244 PE=3 SV=1
CCG020923.1	415	4	25	0.41416612	2.105920287	2.346169394	Up	0.0005384	0.00084414	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1

CCG015479.1	2796	4	25	0.06147315	0.312574005	2.346169394	Up	0.0005384	0.00084419	AAEL006287-PA OS=Aedes aegypti GN=PIWI7 PE=4 SV=1
CCG019909.1	988	4	25	0.17396654	0.884571781	2.346169394	Up	0.0005384	0.00084398	AAEL000135-PA OS=Aedes aegypti GN=AAEL000135 PE=4 SV=1
CCG006313.1	1278	4	25	0.13449056	0.683847355	2.346169394	Up	0.0005384	0.00084424	AAEL000346-PA OS=Aedes aegypti GN=AAEL006348 PE=4 SV=1
CCG012289.1	1929	50	25	1.11378265	0.453062167	-1.297686796	Down	0.0001524	0.00024635	AAEL002824-PA OS=Aedes aegypti GN=AAEL002824 PE=4 SV=1
CCG012520.1	1257	50	25	1.70921776	0.695272012	-1.297686796	Down	0.0001524	0.00024634	AAEL000822-PA OS=Aedes aegypti GN=AAEL008822 PE=4 SV=1
CCG021396.1	1323	52	25	1.68890869	0.660587241	-1.354270325	Down	6.54E-05	0.00010866	AAEL001632-PA OS=Aedes aegypti GN=AAEL001632 PE=4 SV=1
CCG004147.1	1095	52	25	2.04057187	0.798134173	-1.354270325	Down	6.54E-05	0.00010867	AAEL007002-PA OS=Aedes aegypti GN=AAEL007002 PE=4 SV=1
CCG011327.1	1662	52	25	1.3444201	0.525846522	-1.354270325	Down	6.54E-05	0.00010866	AAEL005869-PA (Fragment) OS=Aedes aegypti GN=AAEL005869 PE=4 SV=1
CCG003563.1	948	53	25	2.40231639	0.921895485	-1.381751061	Down	4.25E-05	7.16E-05	AAEL001297-PA OS=Aedes aegypti GN=AAEL001297 PE=4 SV=1
CCG027336.1	3497	59	25	0.72496836	0.249916191	-1.536473656	Down	2.88E-06	5.17E-06	AAEL001201-PA OS=Aedes aegypti GN=AAEL001201 PE=4 SV=1
CCG026569.1	773	59	25	3.27970808	1.130604035	-1.536473656	Down	2.88E-06	5.17E-06	-
CCG011883.1	882	60	25	2.9231112	0.990880861	-1.560721202	Down	1.81E-06	3.29E-06	AAEL005633-PA OS=Aedes aegypti GN=AAEL005633 PE=4 SV=1
CCG027050.1	1043	61	25	2.5130909	0.837926097	-1.584567944	Down	1.13E-06	2.07E-06	AAEL007027-PA OS=Aedes aegypti GN=AAEL007027 PE=3 SV=1
CCG028173.1	465	70	25	6.4685622	1.879477246	-1.783113623	Down	1.37E-08	2.72E-08	-
CCG003400.1	856	77	25	3.86526818	1.020977709	-1.920617147	Down	3.63E-10	7.62E-10	-
CCG015144.1	513	78	25	6.53340994	1.703619726	-1.939232825	Down	2.14E-10	4.52E-10	IAP-antagonist michelob_x OS=Aedes albopictus PE=4 SV=1
CCG000451.1	6705	86	25	0.55114052	0.13034406	-2.080095361	Down	2.77E-12	6.51E-12	Putative uncharacterized protein OS=Drosophila simulans GN=pol PE=4 SV=1
CCG028075.1	1152	86	25	3.20781005	0.758643159	-2.080095361	Down	2.77E-12	6.51E-12	AAEL012612-PA OS=Aedes aegypti GN=AAEL012612 PE=4 SV=1

CCG009200.1	621	89	25	6.15830335	1.407338034	-2.129564037	Down	5.21E-13	1.35E-12	AAEL003114-PA OS=Aedes aegypti GN=AAEL003114 PE=4 SV=1
CCG023491.1	741	91	25	5.27698495	1.179429041	-2.161625247	Down	1.69E-13	4.97E-13	AAEL003506-PA OS=Aedes aegypti GN=AAEL003506 PE=3 SV=1
CCG007097.1	621	93	25	6.43508103	1.407338034	-2.192989418	Down	5.43E-14	1.81E-13	AAEL003114-PA OS=Aedes aegypti GN=AAEL003114 PE=4 SV=1
CCG027297.1	558	111	25	8.54774291	1.566231038	-2.448246473	Down	1.40E-18	5.37E-18	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002735 PE=4 SV=1
CCG002137.1	1302	125	25	4.12535855	0.671241874	-2.619614891	Down	2.61E-22	1.04E-21	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009310 PE=4 SV=1
CCG018760.1	900	139	25	6.63643679	0.971063244	-2.772771679	Down	3.80E-26	1.56E-25	AAEL010866-PA OS=Aedes aegypti GN=AAEL010866 PE=3 SV=1
CCG004517.1	2185	172	25	3.38251458	0.399980283	-3.080095361	Down	1.59E-35	6.92E-35	AAEL001687-PA OS=Aedes aegypti GN=AAEL001687 PE=4 SV=1
CCG018218.1	627	182	25	12.4728735	1.393870685	-3.161625247	Down	1.93E-38	8.57E-38	AAEL001156-PA OS=Aedes aegypti GN=AAEL001156 PE=4 SV=1
CCG009774.1	1581	207	25	5.62601838	0.552787425	-3.347317564	Down	7.61E-46	3.49E-45	AAEL006737-PA OS=Aedes aegypti GN=AAEL006737 PE=4 SV=1
CCG023442.1	1611	252	25	6.72152273	0.542493432	-3.63111053	Down	1.66E-59	8.00E-59	AAEL002666-PA OS=Aedes aegypti GN=AAEL002666 PE=4 SV=1
CCG022691.1	594	256	25	18.5189429	1.471307945	-3.653830606	Down	9.77E-61	4.73E-60	AAEL005253-PA OS=Aedes aegypti GN=AAEL005253 PE=4 SV=1
CCG017440.1	1584	406	25	11.0137072	0.551740479	-4.319166524	Down	3.58E-108	1.92E-107	AAEL006241-PA OS=Aedes aegypti GN=AAEL006241 PE=4 SV=1
CCG016379.1	240	0	26	0.001	3.78714665	11.88689557	Up	2.09E-07	3.96E-07	AAEL014317-PA OS=Aedes aegypti GN=AAEL014317 PE=4 SV=1
CCG023082.1	588	0	26	0.001	1.545774143	10.59411382	Up	2.09E-07	3.96E-07	AAEL012787-PA OS=Aedes aegypti GN=AAEL012787 PE=4 SV=1
CCG020796.1	954	0	26	0.001	0.952741296	9.895940712	Up	2.09E-07	3.96E-07	AGAP000537-PA OS=Anopheles gambiae GN=AgaP_AGAP000537 PE=4 SV=5
CCG007077.1	1783	0	26	0.001	0.509767356	8.993695181	Up	2.09E-07	3.96E-07	Polyprotein OS=Antheraea mylitta PE=4 SV=1

CCG013467.1	426	1	26	0.10086792	2.133603747	4.402752922	Up	0.0005897	0.00091838	-
CCG011594.1	1560	2	26	0.0550894	0.582637946	3.402752922	Up	0.0001122	0.00018266	AAEL010194-PA OS=Aedes aegypti GN=AAEL010194 PE=4 SV=1
CCG003349.1	942	2	26	0.09123086	0.964878127	3.402752922	Up	0.0001122	0.00018265	AAEL005554-PA OS=Aedes aegypti GN=AAEL005554 PE=4 SV=1
CCG012401.1	1655	2	26	0.05192717	0.549193472	3.402752922	Up	0.0001122	0.00018268	Putative uncharacterized protein (Fragment) OS=Camponotus floridanus GN=EAG_00896 PE=4 SV=1
CCG011245.1	657	2	26	0.13080589	1.383432566	3.402752922	Up	0.0001122	0.00018267	AAEL010636-PA OS=Aedes aegypti GN=AAEL010636 PE=4 SV=1
CCG004282.1	396	4	26	0.43403772	2.295240394	2.402752922	Up	0.0003495	0.00055383	AAEL012397-PA OS=Aedes aegypti GN=AAEL012397 PE=4 SV=1
CCG009777.1	3455	4	26	0.04974788	0.263072416	2.402752922	Up	0.0003495	0.0005539	Reverse transcriptase (Protease) (Fragment) OS=Anopheles stephensi PE=4 SV=1
CCG025020.1	354	4	26	0.48553372	2.567557051	2.402752922	Up	0.0003495	0.00055387	-
CCG006962.1	1295	4	26	0.13272505	0.701865016	2.402752922	Up	0.0003495	0.00055394	-
CCG002300.1	1635	4	26	0.10512473	0.555911435	2.402752922	Up	0.0003495	0.0005538	AAEL009771-PA OS=Aedes aegypti GN=AAEL009771 PE=3 SV=1
CCG025721.1	712	4	26	0.241403	1.276566287	2.402752922	Up	0.0003495	0.00055376	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001628 PE=4 SV=1
CCG004786.1	288	51	26	7.60922384	3.155955542	-1.26967242	Down	0.0001695	0.00027328	AAEL012922-PA OS=Aedes aegypti GN=AAEL012922 PE=4 SV=1
CCG009956.1	963	54	26	2.40951783	0.943837171	-1.35213458	Down	4.82E-05	8.10E-05	AAEL008651-PA OS=Aedes aegypti GN=AAEL008651 PE=4 SV=1
CCG017999.1	1268	57	26	1.93160479	0.716810092	-1.430137092	Down	1.31E-05	2.28E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018804 PE=4 SV=1
CCG012579.1	1575	62	26	1.69150701	0.577089013	-1.551443388	Down	1.35E-06	2.47E-06	AAEL002031-PA OS=Aedes aegypti GN=CYP12F7 PE=3 SV=2
CCG018373.1	312	66	26	9.08975155	2.913189731	-1.641641197	Down	2.03E-07	3.84E-07	-
CCG028114.1	1581	99	26	2.69070444	0.574898922	-2.226603698	Down	4.62E-15	1.67E-14	AAEL006737-PA OS=Aedes aegypti GN=AAEL006737 PE=4 SV=1

CCG014909.1	1362	109	26	3.43884073	0.667338617	-2.365431403	Down	1.35E-17	5.15E-17	AAEL006350-PA OS=Aedes aegypti GN=AAEL006350 PE=4 SV=1
CCG027236.1	1515	116	26	3.29009189	0.599944024	-2.455228073	Down	2.06E-19	7.97E-19	AAEL007684-PA OS=Aedes aegypti GN=AAEL007684 PE=4 SV=1
CCG002545.1	1941	135	26	2.98862142	0.468271611	-2.674062675	Down	1.68E-24	6.79E-24	AAEL012947-PA OS=Aedes aegypti GN=AAEL012947 PE=4 SV=1
CCG021107.1	681	155	26	9.78018923	1.334677234	-2.873371483	Down	4.71E-30	1.98E-29	AAEL002305-PA OS=Aedes aegypti GN=AAEL002305 PE=4 SV=1
CCG001408.1	2055	182	26	3.80559207	0.442294499	-3.105041718	Down	8.61E-38	3.80E-37	AAEL012947-PA OS=Aedes aegypti GN=AAEL012947 PE=4 SV=1
CCG013006.1	2019	256	26	5.44836655	0.45018088	-3.597247078	Down	5.88E-60	2.83E-59	AAEL011779-PA OS=Aedes aegypti GN=AAEL011779 PE=4 SV=1
CCG009020.1	2082	354	26	7.3060932	0.436558692	-4.064852628	Down	1.36E-90	7.08E-90	AAEL003135-PA OS=Aedes aegypti GN=AAEL003135 PE=4 SV=1
CCG002135.1	1779	477	26	11.521396	0.510913545	-4.495092534	Down	3.62E-130	2.01E-129	AAEL001835-PA OS=Aedes aegypti GN=AAEL001835 PE=3 SV=1
CCG020789.1	909	1191	26	56.3002793	0.999906706	-5.815204776	Down	0	0	AAEL008092-PA OS=Aedes aegypti GN=AAEL008092 PE=4 SV=1
CCG020663.1	663	1243	26	80.560151	1.370912815	-5.876857659	Down	0	0	AGAP003308-PA OS=Anopheles gambiae GN=AgaP_AGAP003308 PE=4 SV=5
CCG001710.1	735	0	27	0.001	1.284181596	10.32663351	Up	1.15E-07	2.21E-07	LINE-1 retrotransposon-like element, putative (Fragment) OS=Aspergillus flavus (strain ATCC 200026 / FGSC A1120 / NRRL 3357 / JCM 12722 / SRRCL167) GN=AFLA_063980 PE=4 SV=1
CCG015990.1	741	0	27	0.001	1.273783364	10.31490422	Up	1.15E-07	2.21E-07	Hypsin-2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000771 PE=3 SV=1
CCG006026.1	855	0	27	0.001	1.103945582	10.10845334	Up	1.15E-07	2.21E-07	AAEL014239-PA OS=Aedes aegypti GN=AAEL014239 PE=4 SV=1
CCG006931.1	942	0	27	0.001	1.001988825	9.968650703	Up	1.15E-07	2.21E-07	AAEL008202-PA OS=Aedes aegypti GN=AAEL008202 PE=3 SV=1
CCG007668.1	1260	0	27	0.001	0.749105931	9.549025934	Up	1.15E-07	2.21E-07	-

CCG012578.1	1808	0	27	0.001	0.522053912	9.02805499	Up	1.15E-07	2.21E-07	AAEL001960-PA OS=Aedes aegypti GN=AAEL001960 PE=3 SV=1
CCG024631.1	519	1	27	0.08279332	1.818638676	4.457200706	Up	0.0005885	0.00091666	AAEL014348-PB OS=Aedes aegypti GN=CASPS8 PE=4 SV=1
CCG003047.1	324	1	27	0.13262264	2.913189731	4.457200706	Up	0.0005885	0.00091666	-
CCG018098.1	1053	2	27	0.08161393	0.896366071	3.457200706	Up	0.0001045	0.00017044	AAEL011917-PA OS=Aedes aegypti GN=AAEL011917 PE=3 SV=1
CCG019808.1	859	3	27	0.15006892	1.098804974	2.872238205	Up	0.0001061	0.00017287	Zinc knuckle domain protein (Fragment) OS=Wolbachia endosymbiont of Drosophila ananassae GN=WwAna1051 PE=4 SV=1
CCG026255.1	1727	4	27	0.09952457	0.546539359	2.457200706	Up	0.0002299	0.00036822	-
CCG001934.1	3426	4	27	0.05016898	0.275503057	2.457200706	Up	0.0002299	0.0003682	AAEL002146-PA OS=Aedes aegypti GN=AAEL002146 PE=4 SV=1
CCG013608.1	1236	5	27	0.17382579	0.763651677	2.135272611	Up	0.0006384	0.00099182	AAEL010207-PA OS=Aedes aegypti GN=AAEL010207 PE=3 SV=1
CCG021236.1	1182	5	27	0.18176707	0.798539317	2.135272611	Up	0.0006384	0.0009917	AAEL005444-PA OS=Aedes aegypti GN=CASPS19 PE=3 SV=1
CCG021063.1	498	5	27	0.43142304	1.895328259	2.135272611	Up	0.0006384	0.00099176	AAEL005128-PA OS=Aedes aegypti GN=AAEL005128 PE=4 SV=1
CCG017352.1	686	50	27	3.13190486	1.375908853	-1.186655484	Down	0.000416	0.00065705	AAEL007131-PA OS=Aedes aegypti GN=AAEL007131 PE=4 SV=1
CCG011493.1	306	54	27	7.58289434	3.084553833	-1.297686796	Down	8.26E-05	0.00013624	AAEL006079-PA OS=Aedes aegypti GN=AAEL006079 PE=4 SV=1
CCG015926.1	1821	56	27	1.32141963	0.518327003	-1.350154216	Down	3.56E-05	6.02E-05	CYP302a1 OS=Aedes aegypti PE=2 SV=1
CCG010087.1	888	58	27	2.80658177	1.062920578	-1.400780289	Down	1.50E-05	2.61E-05	AAEL008909-PA OS=Aedes aegypti GN=AAEL008909 PE=4 SV=1
CCG018039.1	897	62	27	2.9700374	1.052255823	-1.496995604	Down	2.53E-06	4.55E-06	AAEL010395-PA OS=Aedes aegypti GN=AAEL010395 PE=4 SV=1
CCG018933.1	707	66	27	4.01131893	1.335040273	-1.587193413	Down	3.96E-07	7.41E-07	Putative gag-pol polyprotein OS=Oryza sativa subsp. japonica GN=OSJNBa0013O08.17 PE=4 SV=1

CCG028223.1	703	68	27	4.15638969	1.342636519	-1.630262135	Down	1.53E-07	2.90E-07	Copia OS=Drosophila simulans GN=GIP PE=4 SV=1
CCG021421.1	676	73	27	4.64022282	1.396262534	-1.732623853	Down	1.33E-08	2.64E-08	AAEL009215-PA OS=Aedes aegypti GN=AAEL009215 PE=4 SV=1
CCG001518.1	2120	76	27	1.54042445	0.445223336	-1.790726807	Down	2.94E-09	5.98E-09	AAEL010599-PA OS=Aedes aegypti GN=AAEL010599 PE=4 SV=1
CCG024807.1	1592	78	27	2.10530107	0.592885347	-1.828201513	Down	1.06E-09	2.18E-09	Gag-like protein OS=Aedes aegypti PE=4 SV=1
CCG026376.1	721	80	27	4.76779302	1.309117161	-1.864727389	Down	3.76E-10	7.86E-10	-
CCG016011.1	1131	82	27	3.11540074	0.834547721	-1.900351299	Down	1.32E-10	2.82E-10	AAEL002730-PA OS=Aedes aegypti GN=SRPN21 PE=3 SV=1
CCG018055.1	1500	89	27	2.54953759	0.629248982	-2.018532725	Down	3.13E-12	7.32E-12	AAEL011332-PA OS=Aedes aegypti GN=AAEL011332 PE=4 SV=1
CCG024092.1	780	97	27	5.34367212	1.210094196	-2.142712136	Down	3.74E-14	1.29E-13	AAEL009876-PA OS=Aedes aegypti GN=AAEL009876 PE=3 SV=1
CCG022142.1	2031	100	27	2.11569348	0.464733369	-2.186655484	Down	6.87E-15	2.47E-14	-
CCG023656.1	1581	105	27	2.85377744	0.597010419	-2.257044812	Down	3.91E-16	1.46E-15	AAEL012404-PA OS=Aedes aegypti GN=AAEL012404 PE=4 SV=1
CCG026389.1	1719	117	27	2.92464162	0.549082881	-2.413164014	Down	3.35E-19	1.30E-18	AAEL002741-PA OS=Aedes aegypti GN=SCRB6 PE=4 SV=2
CCG027640.1	1359	118	27	3.73099977	0.6945353	-2.425442343	Down	1.84E-19	7.15E-19	AAEL012720-PA OS=Aedes aegypti GN=AAEL012720 PE=4 SV=1
CCG025140.1	1239	132	27	4.5778894	0.761802642	-2.587193413	Down	3.62E-23	1.45E-22	AAEL008745-PA OS=Aedes aegypti GN=AAEL008745 PE=4 SV=1
CCG015397.1	1056	141	27	5.73743616	0.893819577	-2.682350646	Down	1.31E-25	5.37E-25	AAEL004295-PA OS=Aedes aegypti GN=AAEL004295 PE=4 SV=1
CCG027434.1	345	143	27	17.8106436	2.735865139	-2.702670631	Down	3.73E-26	1.53E-25	AAEL014852-PA OS=Aedes aegypti GN=AAEL014852 PE=2 SV=1
CCG027973.1	912	146	27	6.87892681	1.034948983	-2.732623853	Down	5.58E-27	2.30E-26	AAEL013483-PA OS=Aedes aegypti GN=AAEL013483 PE=4 SV=1
CCG021639.1	886	146	27	7.08079148	1.065319947	-2.732623853	Down	5.58E-27	2.30E-26	Polyprotein OS=Antheraea mylitta PE=4 SV=1
CCG022133.1	1203	155	27	5.53641635	0.784599728	-2.818923699	Down	1.77E-29	7.44E-29	AGAP000592-PA OS=Anopheles gambiae GN=AgaP_AGAP000592 PE=4 SV=3

CCG021550.1	1851	159	27	3.69107931	0.509926241	-2.855682249	Down	1.34E-30	5.67E-30	AGAP004287-PA OS=Anopheles gambiae GN=AgaP_AGAP004287 PE=4 SV=5
CCG017042.1	1137	169	27	6.38688228	0.830143776	-2.94367873	Down	1.99E-33	8.52E-33	AAEL010232-PA OS=Aedes aegypti GN=AAEL010232 PE=4 SV=1
CCG007838.1	888	176	27	8.51652398	1.062920578	-3.002230913	Down	1.98E-35	8.62E-35	AAEL002791-PA OS=Aedes aegypti GN=AAEL002791 PE=4 SV=1
CCG026670.1	1206	178	27	6.3421333	0.782647987	-3.018532725	Down	5.28E-36	2.30E-35	AAEL014394-PA OS=Aedes aegypti GN=CYP301A1 PE=3 SV=2
CCG001530.1	2259	449	27	8.54068652	0.417828009	-4.353370929	Down	4.32E-120	2.36E-119	AAEL002354-PA OS=Aedes aegypti GN=HPX5 PE=4 SV=2
CCG013463.1	1215	572	27	20.229373	0.776850595	-4.702670631	Down	3.69E-160	2.12E-159	AAEL005530-PA OS=Aedes aegypti GN=AAEL005530 PE=4 SV=1
CCG003930.1	354	0	28	0.001	2.76506144	11.43309582	Up	6.36E-08	1.23E-07	-
CCG003756.1	390	0	28	0.001	2.509824999	11.29337106	Up	6.36E-08	1.23E-07	-
CCG011336.1	558	0	28	0.001	1.754178763	10.77658006	Up	6.36E-08	1.23E-07	AAEL010291-PA OS=Aedes aegypti GN=AAEL010291 PE=4 SV=1
CCG027450.1	669	0	28	0.001	1.463126681	10.51483897	Up	6.36E-08	1.23E-07	-
CCG028048.1	848	0	28	0.001	1.154282724	10.17278092	Up	6.36E-08	1.23E-07	-
CCG027564.1	2505	0	28	0.001	0.390751197	8.610106484	Up	6.36E-08	1.23E-07	Protein translocase subunit SecA OS=Wolbachia endosymbiont of Muscidifurax uniraptor GN=secA PE=3 SV=1
CCG007712.1	1680	2	28	0.05115445	0.582637946	3.509668126	Up	0.0001	0.00016314	Chaperone protein DnaK OS=Wolbachia pipientis wAlbB GN=dnaK PE=3 SV=1
CCG017638.1	588	2	28	0.14615556	1.664679846	3.509668126	Up	0.0001	0.00016313	AAEL011649-PA OS=Aedes aegypti GN=AAEL011649 PE=4 SV=1
CCG006044.1	1140	2	28	0.0753855	0.858624342	3.509668126	Up	0.0001	0.00016311	tenon-g OS=Culex quinquefasciatus GN=CpipJ_CPIJ008961 PE=4 SV=1
CCG024584.1	2261	2	28	0.0380095	0.432919836	3.509668126	Up	0.0001	0.00016312	Pol polyprotein (Fragment) OS=Anopheles gambiae PE=4 SV=1

CCG004237.1	558	3	28	0.23102008	1.754178763	2.924705625	Up	8.51E-05	0.00014033	AAEL010421-PA OS=Aedes aegypti GN=AAEL010421 PE=4 SV=1
CCG023916.1	273	3	28	0.47219489	3.585464284	2.924705625	Up	8.51E-05	0.00014032	-
CCG027873.1	493	4	28	0.34863882	1.985459938	2.509668126	Up	0.0001545	0.00024957	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001628 PE=4 SV=1
CCG027942.1	1104	4	28	0.15568744	0.886622962	2.509668126	Up	0.0001545	0.00024955	RUNX1 OS=Aedes aegypti PE=2 SV=1
CCG025024.1	1229	5	28	0.17481584	0.796445687	2.187740031	Up	0.0004152	0.00065583	-
CCG017915.1	404	51	28	5.42439719	2.422850865	-1.162757216	Down	0.0004521	0.00071238	-
CCG007837.1	255	52	28	8.76245569	3.838555881	-1.190771592	Down	0.0003062	0.00048691	AAEL002812-PA OS=Aedes aegypti GN=AAEL002812 PE=4 SV=1
CCG015890.1	1052	52	28	2.12397928	0.930448431	-1.190771592	Down	0.0003062	0.00048688	Endonuclease-reverse transcriptase OS=Schistosoma mansoni PE=2 SV=1
CCG027002.1	764	54	28	3.03712784	1.28119339	-1.245219376	Down	0.000138	0.00022366	-
CCG014141.1	784	62	28	3.39811677	1.248509885	-1.444528185	Down	4.61E-06	8.21E-06	-
CCG028587.1	429	66	28	6.6107284	2.28165909	-1.534725993	Down	7.53E-07	1.39E-06	AAEL011232-PA OS=Aedes aegypti GN=AAEL011232 PE=4 SV=1
CCG022041.1	1098	68	28	2.66114932	0.891467896	-1.577794715	Down	2.97E-07	5.57E-07	AAEL006265-PA (Fragment) OS=Aedes aegypti GN=AAEL006265 PE=4 SV=1
CCG026019.1	585	70	28	5.14167765	1.673216666	-1.619614891	Down	1.15E-07	2.21E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009687 PE=4 SV=1
CCG007985.1	1680	70	28	1.79040561	0.582637946	-1.619614891	Down	1.15E-07	2.21E-07	AAEL004991-PA OS=Aedes aegypti GN=AAEL004991 PE=4 SV=1
CCG009094.1	491	72	28	6.30106088	1.993547352	-1.660256876	Down	4.40E-08	8.57E-08	-
CCG013828.1	1425	73	28	2.20125658	0.686899473	-1.680156433	Down	2.71E-08	5.32E-08	AAEL011874-PA OS=Aedes aegypti GN=AAEL011874 PE=4 SV=1
CCG026250.1	1129	74	28	2.81643965	0.866990035	-1.69978524	Down	1.66E-08	3.28E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009588 PE=4 SV=1
CCG023439.1	809	75	28	3.98359715	1.209927997	-1.719150565	Down	1.01E-08	2.02E-08	-

CCG025011.1	525	77	28	6.30222774	1.864441428	-1.757118415	Down	3.74E-09	7.56E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012798 PE=4 SV=1
CCG017698.1	1275	100	28	3.37017526	0.767711176	-2.134188064	Down	1.76E-14	6.19E-14	AAEL013648-PA OS=Aedes aegypti GN=AAEL013648 PE=4 SV=1
CCG007852.1	1233	109	28	3.79862212	0.793861922	-2.258516199	Down	1.05E-16	3.94E-16	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010318 PE=4 SV=1
CCG024729.1	885	119	28	5.77785132	1.106024576	-2.385149637	Down	2.97E-19	1.15E-18	AAEL010943-PA OS=Aedes aegypti GN=AAEL010943 PE=4 SV=1
CCG011721.1	1290	126	28	4.19704385	0.758784302	-2.467611798	Down	4.46E-21	1.76E-20	AAEL013637-PA OS=Aedes aegypti GN=AAEL013637 PE=4 SV=1
CCG017541.1	399	169	28	18.2002134	2.453212405	-2.89121131	Down	7.79E-33	3.33E-32	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002607 PE=4 SV=1
CCG015384.1	558	176	28	13.5531779	1.754178763	-2.949763493	Down	8.05E-35	3.48E-34	pHCl, isoform 1 OS=Drosophila melanogaster GN=pHCl PE=4 SV=1
CCG015896.1	1389	176	28	5.44468919	0.704702484	-2.949763493	Down	8.05E-35	3.48E-34	AAEL006700-PA OS=Aedes aegypti GN=AAEL006700 PE=3 SV=1
CCG027202.1	2097	182	28	3.72937134	0.466777182	-2.998126514	Down	1.55E-36	6.78E-36	AAEL011512-PA OS=Aedes aegypti GN=AAEL011512 PE=4 SV=1
CCG024949.1	627	183	28	12.5414058	1.561135167	-3.006031712	Down	8.01E-37	3.51E-36	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015249 PE=4 SV=1
CCG008370.1	807	199	28	10.5960064	1.212926579	-3.126956495	Down	1.87E-41	8.41E-41	AAEL005008-PA OS=Aedes aegypti GN=AAEL005008 PE=2 SV=1
CCG013417.1	1440	212	28	6.32609982	0.679744271	-3.218252329	Down	2.88E-45	1.32E-44	AAEL007218-PA OS=Aedes aegypti GN=AAEL007218 PE=4 SV=1
CCG025827.1	1710	274	28	6.88520894	0.572416228	-3.588363957	Down	6.27E-64	3.05E-63	AAEL010856-PA OS=Aedes aegypti GN=AAEL010856 PE=4 SV=1
CCG002864.1	1770	460	28	11.1672757	0.553012288	-4.335821925	Down	1.16E-122	6.37E-122	AAEL008603-PA OS=Aedes aegypti GN=AAEL008603 PE=4 SV=1
CCG027885.1	428	0	29	0.001	2.368668286	11.20986046	Up	3.51E-08	6.86E-08	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1

CCG027673.1	428	0	29	0.001	2.368668286	11.20986046	Up	3.51E-08	6.85E-08	AAEL010609-PA OS=Aedes aegypti GN=AAEL010609 PE=4 SV=1
CCG000320.1	816	0	29	0.001	1.242389738	10.2789021	Up	3.51E-08	6.85E-08	AAEL005607-PA OS=Aedes aegypti GN=AAEL005607 PE=3 SV=1
CCG018585.3	1455	0	29	0.001	0.696762905	9.444524007	Up	3.51E-08	6.85E-08	AAEL005905-PA OS=Aedes aegypti GN=CASPS15 PE=3 SV=1
CCG018071.1	1569	0	29	0.001	0.646137684	9.335697808	Up	3.51E-08	6.85E-08	AGAP001625-PA OS=Anopheles gambiae GN=AgaP_AGAP001625 PE=4 SV=5
CCG000275.1	762	1	29	0.05639073	1.330433106	4.560294199	Up	0.0005875	0.00091512	-
CCG014797.1	1152	1	29	0.03730012	0.880026065	4.560294199	Up	0.0005875	0.00091518	AAEL008459-PA OS=Aedes aegypti GN=AAEL008459 PE=4 SV=1
CCG019376.1	612	1	29	0.07021198	1.656519651	4.560294199	Up	0.0005875	0.00091507	AAEL006639-PA OS=Aedes aegypti GN=AAEL006639 PE=4 SV=1
CCG020649.1	643	2	29	0.13365392	1.57665634	3.560294199	Up	9.73E-05	0.00015886	-
CCG001033.1	642	2	29	0.1338621	1.579112191	3.560294199	Up	9.73E-05	0.00015887	-
CCG013766.1	1635	2	29	0.05256237	0.620055062	3.560294199	Up	9.73E-05	0.00015885	AAEL011341-PA OS=Aedes aegypti GN=AAEL011341 PE=3 SV=1
CCG014413.1	418	2	29	0.20559682	2.425334991	3.560294199	Up	9.73E-05	0.00015888	-
CCG018635.1	198	3	29	0.65105659	5.120151648	2.975331698	Up	7.23E-05	0.00011982	-
CCG018713.1	1983	3	29	0.06500716	0.511240558	2.975331698	Up	7.23E-05	0.00011981	AAEL000801-PA OS=Aedes aegypti GN=AAEL000801 PE=3 SV=1
CCG025595.1	2079	4	29	0.08267385	0.48763349	2.560294199	Up	0.0001072	0.00017463	AAEL013981-PA OS=Aedes aegypti GN=AAEL013981 PE=4 SV=1
CCG023310.1	1134	4	29	0.15156873	0.893994732	2.560294199	Up	0.0001072	0.00017464	AAEL009915-PA (Fragment) OS=Aedes aegypti GN=AAEL009915 PE=4 SV=1
CCG009499.1	1586	5	29	0.13546575	0.63921187	2.238366104	Up	0.0002709	0.00043227	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016425 PE=4 SV=1
CCG007414.1	1107	54	29	2.09608462	0.915799482	-1.194593303	Down	0.0002255	0.00036142	AAEL011573-PA OS=Aedes aegypti GN=AAEL011573 PE=4 SV=1
CCG005964.1	2058	55	29	1.14836511	0.492609342	-1.221065515	Down	0.0001518	0.00024549	AAEL009939-PA OS=Aedes aegypti GN=AAEL009939 PE=4 SV=1

CCG019371.1	501	59	29	5.06030807	2.023532987	-1.32234885	Down	2.96E-05	5.02E-05	AAEL005622-PA OS=Aedes aegypti GN=AAEL005622 PE=4 SV=1
CCG006968.1	2541	61	29	1.0315442	0.398972856	-1.370443139	Down	1.27E-05	2.20E-05	Putative uncharacterized protein OS=Drosophila simulans GN=pol PE=4 SV=1
CCG019921.1	878	71	29	3.47477353	1.154658344	-1.589452921	Down	1.39E-07	2.65E-07	AAEL010891-PA OS=Aedes aegypti GN=AAEL010891 PE=4 SV=1
CCG017003.1	4608	73	29	0.68072713	0.220006516	-1.62953036	Down	5.38E-08	1.05E-07	AAEL011867-PA OS=Aedes aegypti GN=AAEL011867 PE=4 SV=1
CCG005768.1	501	74	29	6.34682707	2.023532987	-1.649159167	Down	3.33E-08	6.51E-08	AAEL012131-PA OS=Aedes aegypti GN=AAEL012131 PE=4 SV=1
CCG014242.1	312	96	29	13.2214568	3.249327008	-2.024668302	Down	3.95E-13	1.05E-12	-
CCG016073.1	1173	100	29	3.66323398	0.864271122	-2.083561991	Down	4.40E-14	1.49E-13	Carboxypeptidase B OS=Aedes aegypti GN=CPB-V PE=2 SV=1
CCG002366.1	618	108	29	7.50927401	1.640436936	-2.194593303	Down	4.94E-16	1.81E-15	AAEL009089-PA OS=Aedes aegypti GN=AAEL009089 PE=4 SV=1
CCG013453.1	772	125	29	6.95753475	1.313199516	-2.405490086	Down	2.42E-20	9.49E-20	AAEL008488-PA OS=Aedes aegypti GN=AAEL008488 PE=4 SV=1
CCG016586.1	1161	167	29	6.18083177	0.873204157	-2.823410094	Down	1.08E-31	4.58E-31	Alcohol dehydrogenase z OS=Culex quinquefasciatus GN=CpipJ_CPIJ016721 PE=4 SV=1
CCG017191.1	1236	176	29	6.11866771	0.820218468	-2.89913742	Down	3.17E-34	1.37E-33	-
CCG007946.1	1412	191	29	5.81247827	0.717981605	-3.017134629	Down	1.67E-38	7.40E-38	AAEL011928-PA OS=Aedes aegypti GN=AAEL011928 PE=4 SV=1
CCG021522.1	648	275	29	18.2356127	1.564490781	-3.542993609	Down	1.80E-63	8.77E-63	AAEL000609-PA OS=Aedes aegypti GN=AAEL000609 PE=4 SV=1
CCG028648.1	720	346	29	20.6493447	1.408041703	-3.874334029	Down	1.91E-85	9.81E-85	AAEL002780-PA OS=Aedes aegypti GN=AAEL002780 PE=4 SV=1
CCG008951.2	1860	472	29	10.9041477	0.545048401	-4.32234885	Down	1.48E-125	8.16E-125	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001238 PE=4 SV=1
CCG025962.1	3201	2197	29	29.4921921	0.316710411	-6.541024955	Down	0	0	AAEL004897-PA OS=Aedes aegypti GN=AAEL004897 PE=4 SV=1

CCG022421.1	3831	37727	29	423.15823	0.264628041	-10.64301556	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_08282 PE=4 SV=1
CCG023174.1	1140	0	30	0.001	0.919954652	9.845418937	Up	1.93E-08	3.82E-08	Yellow-g OS=Culex quinquefasciatus GN=CpipJ_CPIJ008961 PE=4 SV=1
CCG006327.1	537	1	30	0.08001813	1.952976356	4.609203799	Up	0.0005873	0.00091501	AAEL007972-PA OS=Aedes aegypti GN=AAEL007972 PE=3 SV=1
CCG025091.1	1446	1	30	0.02971628	0.725275452	4.609203799	Up	0.0005873	0.0009149	AAEL012230-PA OS=Aedes aegypti GN=AAEL012230 PE=4 SV=1
CCG016472.1	873	1	30	0.04922077	1.201315353	4.609203799	Up	0.0005873	0.00091495	AAEL004584-PA OS=Aedes aegypti GN=AAEL004584 PE=4 SV=1
CCG012380.1	966	2	30	0.08896425	1.085660769	3.609203799	Up	9.58E-05	0.00015639	AAEL014506-PA (Fragment) OS=Aedes aegypti GN=AAEL014506 PE=4 SV=1
CCG020165.1	537	2	30	0.16003626	1.952976356	3.609203799	Up	9.58E-05	0.00015638	AAEL006623-PA OS=Aedes aegypti GN=AAEL006623 PE=4 SV=1
CCG020232.2	777	4	30	0.22120841	1.349740416	2.609203799	Up	7.76E-05	0.00012835	Uncharacterized protein OS=Anopheles darlingi GN=AND_02384 PE=4 SV=1
CCG001244.1	942	4	30	0.18246172	1.113320916	2.609203799	Up	7.76E-05	0.00012834	AAEL005554-PA OS=Aedes aegypti GN=AAEL005554 PE=4 SV=1
CCG005434.1	484	4	30	0.35512177	2.166835337	2.609203799	Up	7.76E-05	0.00012833	-
CCG017071.1	1341	5	30	0.16021527	0.782064357	2.287275705	Up	0.0001781	0.00028698	AAEL009669-PA OS=Aedes aegypti GN=AAEL009669 PE=4 SV=1
CCG005622.1	2325	6	30	0.11088964	0.451074539	2.024241299	Up	0.0004776	0.00075159	Translation initiation factor IF-2 OS=Wolbachia endosymbiont of Muscidifurax uniraptor GN=infB PE=3 SV=1
CCG000867.1	673	57	30	3.63933859	1.55831843	-1.223686215	Down	0.0001119	0.00018216	AAEL000532-PA OS=Aedes aegypti GN=AAEL000532 PE=4 SV=1
CCG026655.1	1329	58	30	1.87527811	0.789125887	-1.248777196	Down	7.49E-05	0.00012405	AAEL009943-PA OS=Aedes aegypti GN=AAEL009943 PE=4 SV=1
CCG012313.1	804	62	30	3.3135865	1.304413312	-1.344992511	Down	1.43E-05	2.49E-05	GK19301 OS=Drosophila willistoni GN=GK19301 PE=4 SV=1

CCG006788.1	369	68	30	7.91854188	2.842136323	-1.478259042	Down	1.04E-06	1.92E-06	AAEL010909-PA OS=Aedes aegypti GN=AAEL010909 PE=4 SV=1
CCG028473.1	2265	74	30	1.40386771	0.463023533	-1.600249566	Down	6.52E-08	1.26E-07	Ribbon OS=Culex quinquefasciatus GN=CripJ_CPIJ004725 PE=4 SV=1
CCG003896.1	432	77	30	7.65895733	2.427658109	-1.657582741	Down	1.55E-08	3.07E-08	AAEL014559-PA OS=Aedes aegypti GN=AAEL014559 PE=4 SV=1
CCG023695.1	1095	78	30	3.06085781	0.957761007	-1.676198419	Down	9.56E-09	1.91E-08	AAEL006696-PA OS=Aedes aegypti GN=AAEL006696 PE=3 SV=1
CCG014313.1	1488	84	30	2.42571082	0.704803967	-1.783113623	Down	4.87E-10	1.01E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004725 PE=4 SV=1
CCG023861.1	348	85	30	10.4954812	3.013644549	-1.800187137	Down	2.94E-10	6.17E-10	Uncharacterized protein OS=Anopheles darlingi GN=AND_09992 PE=3 SV=1
CCG026650.1	623	100	30	6.89722867	1.683384114	-2.03465239	Down	1.07E-13	3.34E-13	-
CCG019646.1	1880	103	30	2.35419291	0.557844842	-2.077296728	Down	2.06E-14	7.23E-14	AGAP005356-PA OS=Anopheles gambiae GN=AGAP005356 PE=4 SV=1
CCG006146.1	452	106	30	10.0769732	2.320239609	-2.118716655	Down	3.90E-15	1.41E-14	-
CCG004915.1	2007	119	30	2.54778197	0.522545243	-2.285613964	Down	2.36E-18	9.04E-18	AAEL014044-PA OS=Aedes aegypti GN=AAEL014044 PE=4 SV=1
CCG006110.1	1155	145	30	5.39446885	0.908007189	-2.570705291	Down	3.95E-25	1.61E-24	AAEL013864-PA OS=Aedes aegypti GN=AAEL013864 PE=4 SV=1
CCG008124.1	1164	177	30	6.53405758	0.900986515	-2.858401751	Down	6.35E-34	2.73E-33	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005093 PE=4 SV=1
CCG026707.1	1284	193	30	6.4588464	0.816782168	-2.983253238	Down	1.84E-38	8.13E-38	AAEL013109-PA OS=Aedes aegypti GN=AAEL013109 PE=4 SV=1
CCG016529.1	1044	242	30	9.96041741	1.004548183	-3.309659438	Down	8.60E-53	4.06E-52	AAEL008496-PA OS=Aedes aegypti GN=AAEL008496 PE=4 SV=1
CCG005396.1	1230	333	30	11.6332696	0.852640897	-3.770174568	Down	1.49E-80	7.58E-80	AAEL010597-PA OS=Aedes aegypti GN=AAEL010597 PE=3 SV=1
CCG007817.1	924	534	30	24.8331583	1.135008986	-4.451492132	Down	1.28E-144	7.21E-144	AAEL012245-PA OS=Aedes aegypti GN=AAEL012245 PE=4 SV=1

CCG010581.1	459	0	31	0.001	2.361016514	11.20519242	Up	1.07E-08	2.13E-08	AAEL007592-PA OS=Aedes aegypti GN=AAEL007592 PE=4 SV=1
CCG004919.1	825	0	31	0.001	1.313583733	10.35929245	Up	1.07E-08	2.13E-08	AAEL002196-PA OS=Aedes aegypti GN=AAEL002196 PE=4 SV=1
CCG024005.1	1296	0	31	0.001	0.836193349	9.707692758	Up	1.07E-08	2.13E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005433 PE=4 SV=1
CCG006728.1	1506	1	31	0.02853236	0.719592683	4.656509514	Up	0.0005871	0.00091487	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006292 PE=4 SV=1
CCG022573.1	417	1	31	0.10304493	2.598816738	4.656509514	Up	0.0005871	0.00091493	AAEL001539-PA (Fragment) OS=Aedes aegypti GN=AAEL001539 PE=4 SV=1
CCG010828.1	447	2	31	0.19225832	2.424399508	3.656509514	Up	9.48E-05	0.0001549	AAEL001249-PA OS=Aedes aegypti GN=AAEL001249 PE=4 SV=1
CCG010000.1	552	2	31	0.15568744	1.963236558	3.656509514	Up	9.48E-05	0.00015492	AAEL009184-PA OS=Aedes aegypti GN=AAEL009184 PE=4 SV=1
CCG013140.1	192	2	31	0.4476014	5.644305104	3.656509514	Up	9.48E-05	0.00015493	Gambicin OS=Aedes triseriatus PE=4 SV=1
CCG000074.1	1434	2	31	0.05992989	0.755722859	3.656509514	Up	9.48E-05	0.00015491	-
CCG010385.1	3978	3	31	0.03240553	0.272424982	3.071547013	Up	5.99E-05	9.96E-05	AAEL009849-PA OS=Aedes aegypti GN=AAEL009849 PE=4 SV=1
CCG000796.1	2115	4	31	0.08126664	0.512390818	2.656509514	Up	5.92E-05	9.86E-05	AAEL007648-PA OS=Aedes aegypti GN=AAEL007648 PE=4 SV=1
CCG002369.1	804	4	31	0.21377977	1.347893756	2.656509514	Up	5.92E-05	9.86E-05	AAEL007626-PA OS=Aedes aegypti GN=GNBPA1 PE=4 SV=2
CCG006024.1	2122	5	31	0.1012482	0.510700556	2.334581419	Up	0.0001187	0.00019301	Putative uncharacterized protein OS=Medicago truncatula GN=MTR_2g061140 PE=4 SV=1
CCG019295.1	980	5	31	0.21923334	1.105823041	2.334581419	Up	0.0001187	0.00019298	-
CCG005659.1	663	5	31	0.32405531	1.634549894	2.334581419	Up	0.0001187	0.00019299	AAEL013352-PA OS=Aedes aegypti GN=AAEL013352 PE=3 SV=1
CCG009410.1	534	5	31	0.40233834	2.029413071	2.334581419	Up	0.0001187	0.000193	-
CCG014092.1	469	6	31	0.54971942	2.310675011	2.071547013	Up	0.0003132	0.00049768	AAEL004644-PA OS=Aedes aegypti GN=AAEL004644 PE=4 SV=1

CCG028547.1	1173	6	31	0.21979404	0.923876027	2.071547013	Up	0.0003132	0.00049761	AAEL009178-PA OS=Aedes aegypti GN=GNBPB4 PE=4 SV=1
CCG004165.1	1752	6	31	0.14715663	0.618553984	2.071547013	Up	0.0003132	0.00049765	AAEL010537-PA OS=Aedes aegypti GN=AAEL010537 PE=4 SV=1
CCG008731.1	1654	6	31	0.1558757	0.655203495	2.071547013	Up	0.0003132	0.00049771	Reverse transcriptase OS=Anopheles gambiae PE=4 SV=1
CCG028077.1	918	64	31	2.99571135	1.180508257	-1.343490486	Down	1.06E-05	1.85E-05	AGAP004169-PA OS=Anopheles gambiae GN=AgaP_AGAP004169 PE=4 SV=4
CCG011806.1	1704	65	31	1.63910373	0.63597804	-1.365858299	Down	6.92E-06	1.22E-05	AAEL005002-PA OS=Aedes aegypti GN=AAEL005002 PE=4 SV=1
CCG014499.1	1362	67	31	2.11378283	0.795672966	-1.409579676	Down	2.92E-06	5.25E-06	AAEL006350-PA OS=Aedes aegypti GN=AAEL006350 PE=4 SV=1
CCG005483.1	272	69	31	10.9004106	3.984215367	-1.452014943	Down	1.21E-06	2.22E-06	-
CCG028443.1	2277	72	31	1.35872679	0.475936135	-1.513415487	Down	3.13E-07	5.88E-07	AAEL014543-PA OS=Aedes aegypti GN=AAEL014543 PE=4 SV=1
CCG027384.1	1989	72	31	1.55546551	0.544849965	-1.513415487	Down	3.13E-07	5.88E-07	AAEL001050-PA OS=Aedes aegypti GN=AAEL001050 PE=4 SV=1
CCG025410.1	1236	77	31	2.67691712	0.876785259	-1.610277026	Down	3.05E-08	6.00E-08	Estradiol 17-beta-dehydrogenase 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ011409 PE=4 SV=1
CCG024929.1	2913	84	31	1.23908606	0.372024229	-1.735807909	Down	1.02E-09	2.10E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003554 PE=4 SV=1
CCG009166.1	813	84	31	4.43967738	1.332972423	-1.735807909	Down	1.02E-09	2.10E-09	AAEL008596-PA OS=Aedes aegypti GN=SPZ3A PE=4 SV=1
CCG014366.1	1195	85	31	3.05642464	0.906867431	-1.752881422	Down	6.19E-10	1.29E-09	-
CCG009242.1	525	111	31	9.0850296	2.064203009	-2.137906352	Down	6.00E-16	2.19E-15	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010048 PE=4 SV=1
CCG017506.1	1113	114	31	4.40121271	0.973680665	-2.1763805	Down	1.11E-16	4.17E-16	AAEL004150-PA OS=Aedes aegypti GN=AAEL004150 PE=4 SV=1
CCG007548.1	1259	114	31	3.89082585	0.860767736	-2.1763805	Down	1.11E-16	4.17E-16	-

CCG005611.1	558	121	31	9.31780984	1.942126487	-2.262353723	Down	2.02E-18	7.74E-18	AAEL001868-PA OS=Aedes aegypti GN=AAEL001868 PE=4 SV=1
CCG024433.1	1434	126	31	3.77558338	0.755722859	-2.320770409	Down	1.10E-19	4.27E-19	AAEL011332-PA OS=Aedes aegypti GN=AAEL011332 PE=4 SV=1
CCG000626.1	1515	151	31	4.28279203	0.715317875	-2.581895225	Down	3.14E-26	1.29E-25	AAEL003516-PA OS=Aedes aegypti GN=AAEL003516 PE=4 SV=1
CCG025179.1	574	153	31	11.4536052	1.887990557	-2.600878328	Down	9.10E-27	3.76E-26	AAEL013387-PA OS=Aedes aegypti GN=AAEL013387 PE=4 SV=1
CCG006666.1	897	203	31	9.72447729	1.208145574	-3.008826403	Down	1.03E-40	4.59E-40	AAEL010866-PA OS=Aedes aegypti GN=AAEL010866 PE=3 SV=1
CCG009432.1	1161	394	31	14.5823217	0.933425133	-3.965542305	Down	8.11E-99	4.29E-98	AAEL013452-PA OS=Aedes aegypti GN=AAEL013452 PE=4 SV=1
CCG025065.1	2145	559	31	11.1981733	0.505224513	-4.470194959	Down	1.01E-151	5.72E-151	AAEL011310-PA OS=Aedes aegypti GN=AAEL011310 PE=4 SV=1
CCG022702.1	3417	912	31	11.4686561	0.317151472	-5.1763805	Down	3.44E-268	2.14E-267	GH13316 OS=Drosophila grimshawi GN=GH13316 PE=4 SV=1
CCG018020.1	1857	2706	31	62.6150252	0.583579203	-6.74543661	Down	0	0	AAEL012090-PA OS=Aedes aegypti GN=AAEL012090 PE=4 SV=1
CCG001623.1	426	0	32	0.001	2.625973842	11.35863683	Up	5.88E-09	1.18E-08	AAEL000262-PA OS=Aedes aegypti GN=AAEL000262 PE=4 SV=1
CCG014440.1	435	0	32	0.001	2.571643349	11.32847486	Up	5.88E-09	1.18E-08	-
CCG021787.1	636	0	32	0.001	1.758907007	10.78046349	Up	5.88E-09	1.18E-08	AAEL009165-PA OS=Aedes aegypti GN=AAEL009165 PE=4 SV=1
CCG014354.1	1112	0	32	0.001	1.005993576	9.974405377	Up	5.88E-09	1.18E-08	U3 small nucleolar ribonucleoprotein component OS=Aedes aegypti PE=2 SV=1
CCG001791.1	1155	0	32	0.001	0.968541001	9.919669314	Up	5.88E-09	1.18E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010663 PE=4 SV=1
CCG023317.1	1275	0	32	0.001	0.877384201	9.777064918	Up	5.88E-09	1.18E-08	AAEL002882-PA OS=Aedes aegypti GN=AAEL002882 PE=4 SV=1
CCG002198.1	561	1	32	0.07659489	1.994055003	4.702313204	Up	0.0005871	0.00091488	AAEL008337-PA OS=Aedes aegypti GN=AAEL008337 PE=4 SV=1

CCG013649.1	2117	2	32	0.04059493	0.528419866	3.702313204	Up	9.43E-05	0.00015411	AAEL010730-PA OS=Aedes aegypti GN=AAEL010730 PE=4 SV=1
CCG026885.1	1275	2	32	0.06740351	0.877384201	3.702313204	Up	9.43E-05	0.00015409	AAEL014814-PA OS=Aedes aegypti GN=AAEL014814 PE=4 SV=1
CCG007709.1	683	2	32	0.12582646	1.637869483	3.702313204	Up	9.43E-05	0.00015408	Uncharacterized protein OS=Anopheles darlingi GN=AND_08768 PE=4 SV=1
CCG028224.1	1078	2	32	0.07972121	1.037722502	3.702313204	Up	9.43E-05	0.0001541	-
CCG025484.1	1205	3	32	0.10697859	0.928352578	3.117350703	Up	5.70E-05	9.50E-05	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG011647.1	651	3	32	0.19801721	1.718379196	3.117350703	Up	5.70E-05	9.50E-05	-
CCG015393.1	1394	4	32	0.1232991	0.80248555	2.702313204	Up	4.78E-05	8.03E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004048 PE=4 SV=1
CCG009487.1	576	4	32	0.29840093	1.942126487	2.702313204	Up	4.78E-05	8.03E-05	-
CCG023152.1	484	5	32	0.44390222	2.311291026	2.380385109	Up	8.08E-05	0.00013332	AAEL006533-PA OS=Aedes aegypti GN=AAEL006533 PE=3 SV=1
CCG003431.1	831	6	32	0.31025079	1.346167096	2.117350703	Up	0.0002055	0.00033027	-
CCG013012.1	966	6	32	0.26689276	1.158038154	2.117350703	Up	0.0002055	0.00033025	AAEL000801-PA OS=Aedes aegypti GN=AAEL000801 PE=3 SV=1
CCG024967.1	813	7	32	0.36997311	1.375971533	1.894958282	Up	0.0005322	0.00083474	AAEL004999-PA OS=Aedes aegypti GN=AAEL004999 PE=4 SV=1
CCG013406.1	1425	60	32	1.80925198	0.78502797	-1.204577392	Down	9.04E-05	0.00014894	AAEL004576-PA OS=Aedes aegypti GN=AAEL004576 PE=4 SV=1
CCG004594.1	699	71	32	4.36459393	1.600378908	-1.447433916	Down	8.98E-07	1.66E-06	AAEL006932-PA OS=Aedes aegypti GN=AAEL006932 PE=4 SV=1
CCG021518.1	1018	72	32	3.03911679	1.098884928	-1.467611798	Down	5.76E-07	1.07E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017997 PE=4 SV=1
CCG009624.2	1528	80	32	2.24972433	0.732110508	-1.619614891	Down	1.43E-08	2.84E-08	-
CCG002432.1	1065	94	32	3.79263385	1.050389537	-1.852275648	Down	1.36E-11	3.01E-11	-
CCG028524.1	2751	115	32	1.79626299	0.406639352	-2.143176847	Down	1.62E-16	6.07E-16	AAEL000025-PA OS=Aedes aegypti GN=AAEL000025 PE=3 SV=1

CCG002970.1	742	122	32	7.06510461	1.507634578	-2.228424134	Down	3.04E-18	1.17E-17	AAEL009705-PA OS=Aedes aegypti GN=AAEL009705 PE=4 SV=1
CCG027792.1	459	141	32	13.1998531	2.437178337	-2.437238149	Down	4.32E-23	1.73E-22	AAEL005291-PA OS=Aedes aegypti GN=AAEL003291 PE=4 SV=1
CCG026373.1	1926	164	32	3.65889744	0.580822875	-2.655238801	Down	3.19E-29	1.34E-28	AAEL006339-PA OS=Aedes aegypti GN=AAEL006339 PE=4 SV=1
CCG027459.1	1323	205	32	6.65819773	0.845551668	-2.977166896	Down	1.12E-40	5.03E-40	AAEL001143-PA OS=Aedes aegypti GN=AAEL001143 PE=4 SV=1
CCG028291.1	744	329	32	19.0014015	1.503581797	-3.65963057	Down	1.05E-77	5.31E-77	AAEL014538-PA OS=Aedes aegypti GN=AAEL014538 PE=4 SV=1
CCG007212.1	1863	1189	32	27.424055	0.600464228	-5.513219796	Down	0	0	AAEL012090-PA OS=Aedes aegypti GN=AAEL012090 PE=4 SV=1
CCG015090.1	2772	1806	32	27.9954332	0.403558751	-6.116268974	Down	0	0	AAEL007483-PA OS=Aedes aegypti GN=AAEL007483 PE=4 SV=1
CCG007494.1	1035	0	33	0.001	1.114611723	10.12232552	Up	3.24E-09	6.60E-09	AAEL000021-PA OS=Aedes aegypti GN=AAEL000021 PE=4 SV=1
CCG005605.1	1086	0	33	0.001	1.062268079	10.05293218	Up	3.24E-09	6.60E-09	AAEL014814-PA OS=Aedes aegypti GN=AAEL014814 PE=4 SV=1
CCG008747.1	1772	0	33	0.001	0.651028856	9.346577681	Up	3.24E-09	6.60E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014613 PE=4 SV=1
CCG012240.1	1478	2	33	0.05814578	0.78052986	3.746707323	Up	9.40E-05	0.00015363	-
CCG016501.1	1509	2	33	0.05695127	0.764495118	3.746707323	Up	9.40E-05	0.00015364	-
CCG002108.1	288	3	33	0.4476014	4.00563588	3.161744822	Up	5.53E-05	9.22E-05	-
CCG016607.1	756	3	33	0.17051482	1.525956526	3.161744822	Up	5.53E-05	9.23E-05	-
CCG021984.1	1071	4	33	0.16048454	1.077145783	2.746707323	Up	4.07E-05	6.87E-05	AAEL007200-PA OS=Aedes aegypti GN=AAEL007200 PE=4 SV=1
CCG011705.1	1103	4	33	0.15582859	1.04589586	2.746707323	Up	4.07E-05	6.87E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007563 PE=4 SV=1
CCG025576.1	1044	5	33	0.20579375	1.105003001	2.424779228	Up	5.67E-05	9.45E-05	-
CCG018538.1	265	5	33	0.81074971	4.353294843	2.424779228	Up	5.67E-05	9.46E-05	-
CCG023878.1	492	5	33	0.43668429	2.344762466	2.424779228	Up	5.67E-05	9.45E-05	GK19301 OS=Drosophila willistoni GN=GK19301 PE=4 SV=1

CCG007181.1	2410	6	33	0.10697859	0.478681798	2.161744822	Up	0.0001354	0.00021952	Putative finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014131 PE=4 SV=1
CCG008143.1	216	6	33	1.19360374	5.34084784	2.161744822	Up	0.0001354	0.00021949	-
CCG022916.1	2772	6	33	0.09300808	0.416169962	2.161744822	Up	0.0001354	0.00021951	AAEL004179-PA OS=Reedes aegypti GN=AAEL004179 PE=4 SV=1
CCG002806.1	1113	6	33	0.23164277	1.036498772	2.161744822	Up	0.0001354	0.00021953	-
CCG017391.1	3627	7	33	0.08293028	0.31806538	1.939352401	Up	0.0003524	0.00055831	AAEL010391-PA OS=Reedes aegypti GN=AAEL010391 PE=4 SV=1
CCG005654.1	661	7	33	0.45505014	1.745269491	1.939352401	Up	0.0003524	0.00055834	-
CCG004086.1	1571	58	33	1.5864065	0.734324082	-1.111273672	Down	0.0003086	0.00049053	AAEL005729-PA OS=Reedes aegypti GN=AAEL005729 PE=4 SV=1
CCG020825.1	2421	64	33	1.13592029	0.476506871	-1.253292677	Down	3.01E-05	5.11E-05	Stopperin OS=Culex quinquefasciatus GN=CpipJ_CPIJ010715 PE=4 SV=1
CCG019099.1	522	67	33	5.51527245	2.210006003	-1.319381867	Down	8.79E-06	1.54E-05	AAEL004964-PA OS=Reedes aegypti GN=AAEL004964 PE=4 SV=1
CCG011995.1	678	71	33	4.49978047	1.701509046	-1.403039796	Down	1.60E-06	2.92E-06	-
CCG021626.1	537	73	33	5.84132333	2.148273992	-1.443117236	Down	6.67E-07	1.24E-06	AAEL004466-PA OS=Reedes aegypti GN=AAEL004466 PE=4 SV=1
CCG015302.1	885	79	33	3.83571642	1.303528964	-1.557073425	Down	4.40E-08	8.58E-08	AAEL011901-PA OS=Reedes aegypti GN=AAEL011901 PE=4 SV=1
CCG020331.1	756	80	33	4.54706186	1.525956526	-1.575220772	Down	2.76E-08	5.43E-08	Galactose-specific G-type lectin OS=Culex quinquefasciatus GN=CpipJ_CPIJ017075 PE=4 SV=1
CCG013640.1	1236	89	33	3.09409901	0.93335205	-1.729026108	Down	3.62E-10	7.60E-10	AAEL009967-PA OS=Reedes aegypti GN=AAEL009967 PE=4 SV=1
CCG008954.1	921	102	33	4.75886312	1.252576692	-1.925718019	Down	4.65E-13	1.21E-12	AAEL010201-PA OS=Reedes aegypti GN=AAEL010201 PE=4 SV=1
CCG021748.1	1542	105	33	2.92595469	0.748134328	-1.967538194	Down	9.44E-14	2.97E-13	GE21538 OS=Drosophila yakuba GN=GE21538 PE=4 SV=1
CCG020258.1	1410	107	33	3.26082383	0.818172435	-1.994759663	Down	3.23E-14	1.12E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001097 PE=4 SV=1

CCG025998.1	3240	115	33	1.52516033	0.356056523	-2.098782728	Down	4.06E-16	1.52E-15	AAEL009780-PA OS=Aedes aegypti GN=AAEL009780 PE=3 SV=1
CCG008761.1	1413	119	33	3.61882408	0.816435339	-2.14811044	Down	4.35E-17	1.65E-16	AAEL001680-PA OS=Aedes aegypti GN=AAEL001680 PE=4 SV=1
CCG020607.1	549	133	33	10.40979	2.101317183	-2.308575112	Down	1.42E-20	5.57E-20	AAEL000455-PA OS=Aedes aegypti GN=AAEL000455 PE=4 SV=1
CCG008910.1	1293	145	33	4.81872507	0.8922066	-2.433201767	Down	1.17E-23	4.68E-23	AAEL005959-PA OS=Aedes aegypti GN=AAEL005959 PE=4 SV=1
CCG015218.1	2220	152	33	2.94207192	0.51965006	-2.50122019	Down	1.70E-25	6.93E-25	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013977 PE=4 SV=1
CCG002479.1	2613	206	33	3.38758719	0.441493737	-2.939793204	Down	2.34E-40	1.05E-39	AAEL000334-PA OS=Aedes aegypti GN=AAEL000334 PE=4 SV=1
CCG005970.1	1076	261	33	10.4229561	1.072140459	-3.281198673	Down	2.29E-56	1.09E-55	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014232 PE=4 SV=1
CCG000177.1	381	349	33	39.360728	3.02788224	-3.700375903	Down	4.41E-83	2.25E-82	AAEL002231-PA OS=Aedes aegypti GN=AAEL002231 PE=4 SV=1
CCG008170.1	915	864	33	40.5747002	1.26079031	-5.008180179	Down	7.94E-250	4.88E-249	AAEL012292-PA OS=Aedes aegypti GN=AAEL012292 PE=4 SV=1
CCG017079.1	954	1530	33	68.9137253	1.209248568	-5.832608614	Down	0	0	AAEL013222-PA OS=Aedes aegypti GN=AAEL013222 PE=4 SV=1
CCG016456.1	627	3775	33	258.709327	1.839909304	-7.135553606	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015249 PE=4 SV=1
CCG019284.1	577	0	34	0.001	2.05993312	11.00838178	Up	1.79E-09	3.66E-09	Histone H2B OS=Aedes aegypti GN=AAEL015682 PE=3 SV=1
CCG004071.1	1275	0	34	0.001	0.932220714	9.86452776	Up	1.79E-09	3.66E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012792 PE=4 SV=1
CCG013348.1	1317	0	34	0.001	0.90249158	9.817769661	Up	1.79E-09	3.66E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005075 PE=4 SV=1
CCG000691.1	594	1	34	0.07233962	2.000978805	4.789776045	Up	0.000587	0.00091495	AAEL003775-PA OS=Aedes aegypti GN=AAEL003775 PE=4 SV=1

CCG019071.1	282	1	34	0.15237495	4.214827696	4.789776045	Up	0.000587	0.00091484	AAEL008863-PA OS=Aedes aegypti GN=AAEL008863 PE=4 SV=1
CCG007916.1	2372	1	34	0.0181154	0.501088284	4.789776045	Up	0.000587	0.00091489	Putative uncharacterized protein OS=Caenorhabditis remanei GN=CRE_08209 PE=4 SV=1
CCG023021.1	490	1	34	0.08769334	2.425676347	4.789776045	Up	0.000587	0.00091501	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG004719.1	4358	2	34	0.01971993	0.272735523	3.789776045	Up	9.38E-05	0.00015339	Putative uncharacterized protein 882 (3' region) (Fragment) OS=Aedes aegypti PE=4 SV=1
CCG011553.1	489	2	34	0.17574534	2.430636831	3.789776045	Up	9.38E-05	0.00015337	AAEL000913-PA OS=Aedes aegypti GN=AAEL000913 PE=4 SV=1
CCG007384.1	1573	2	34	0.05463412	0.755614374	3.789776045	Up	9.38E-05	0.00015338	AAEL006747-PA OS=Aedes aegypti GN=AAEL006747 PE=4 SV=1
CCG016872.1	2110	2	34	0.04072961	0.563308725	3.789776045	Up	9.38E-05	0.00015335	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG020376.1	456	2	34	0.18846375	2.60653818	3.789776045	Up	9.38E-05	0.0001534	-
CCG028479.1	1011	2	34	0.08500442	1.175649268	3.789776045	Up	9.38E-05	0.00015336	AAEL014410-PA OS=Aedes aegypti GN=AAEL014410 PE=4 SV=1
CCG007710.1	1231	4	34	0.13962546	0.965541357	2.789776045	Up	3.64E-05	6.15E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC002657 PE=4 SV=1
CCG018263.1	1713	4	34	0.10033797	0.693859551	2.789776045	Up	3.64E-05	6.15E-05	AAEL009205-PA OS=Aedes aegypti GN=AAEL009205 PE=4 SV=1
CCG018026.1	339	4	34	0.50701752	3.506139853	2.789776045	Up	3.64E-05	6.15E-05	-
CCG024557.1	468	4	34	0.36726269	2.539703868	2.789776045	Up	3.64E-05	6.15E-05	-
CCG004394.1	789	4	34	0.21784403	1.506440317	2.789776045	Up	3.64E-05	6.15E-05	-
CCG008831.1	654	4	34	0.26281183	1.817402768	2.789776045	Up	3.64E-05	6.15E-05	AAEL004416-PA OS=Aedes aegypti GN=AAEL004416 PE=4 SV=1
CCG007628.1	1812	6	34	0.14228389	0.655950006	2.204813544	Up	8.99E-05	0.00014816	AAEL007486-PA (Fragment) OS=Aedes aegypti GN=AAEL007486 PE=4 SV=2

CCG022976.1	327	6	34	0.7884355	3.634805536	2.204813544	Up	8.99E-05	0.00014817	AAEL002527-PA OS=Anopheles aegypti GN=AAEL002527 PE=4 SV=1
CCG021364.1	1325	7	34	0.22700992	0.897042574	1.982421123	Up	0.0002328	0.0003729	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG017799.1	375	8	34	0.91668767	3.169550427	1.789776045	Up	0.0005768	0.00090326	Uncharacterized protein OS=Anopheles darlingi GN=AND_22697 PE=4 SV=1
CCG018606.1	1332	8	34	0.25807648	0.892328386	1.789776045	Up	0.0005768	0.00090332	AAEL001053-PA OS=Anopheles aegypti GN=AAEL001053 PE=4 SV=1
CCG025178.1	1956	60	34	1.31809002	0.607659208	-1.117114551	Down	0.0002282	0.00036562	Putative uncharacterized protein GLEAN_07721 OS=Tribolium castaneum GN=GLEAN_07721 PE=4 SV=1
CCG010439.1	1452	61	34	1.80520235	0.818582238	-1.140961292	Down	0.0001567	0.00025307	AGAP013133-PA OS=Anopheles gambiae GN=AgaP_AGAP013133 PE=4 SV=1
CCG007853.1	1353	61	34	1.93729033	0.8784785	-1.140961292	Down	0.0001567	0.00025306	AAEL014435-PA OS=Anopheles aegypti GN=AAEL014435 PE=4 SV=1
CCG021812.1	561	63	34	4.82547822	2.118683441	-1.187503878	Down	7.28E-05	0.00012058	AAEL012725-PA OS=Anopheles aegypti GN=AAEL012725 PE=4 SV=1
CCG025971.1	1323	65	34	2.11113587	0.898398647	-1.232591768	Down	3.31E-05	5.61E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002877 PE=4 SV=1
CCG006206.1	520	65	34	5.37121683	2.285733481	-1.232591768	Down	3.31E-05	5.61E-05	-
CCG003161.1	2444	66	34	1.16039382	0.486326273	-1.254618074	Down	2.22E-05	3.81E-05	Putative gag-pol protein OS=Drosophila ananassae PE=4 SV=1
CCG008833.1	1146	72	34	2.69966919	1.037156553	-1.380148956	Down	1.83E-06	3.33E-06	AAEL004418-PA OS=Anopheles aegypti GN=AAEL004418 PE=4 SV=1
CCG020168.1	2071	73	34	1.51462609	0.573916664	-1.400048514	Down	1.19E-06	2.18E-06	GG23150 OS=Drosophila erecta GN=GG23150 PE=4 SV=1
CCG016104.1	933	74	34	3.40810328	1.273935059	-1.419677321	Down	7.69E-07	1.42E-06	AAEL009726-PA OS=Anopheles aegypti GN=AAEL009726 PE=3 SV=1

CCG021681.1	1098	83	34	3.24816755	1.082496731	-1.585263386	Down	1.30E-08	2.59E-08	Cerrate determining protein mab2112 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014033 PE=4 SV=1
CCG002883.1	1203	90	34	3.21469336	0.988014472	-1.702077051	Down	4.53E-10	9.44E-10	AAEL007076-PA OS=Aedes aegypti GN=AAEL007878 PE=3 SV=1
CCG021773.1	1479	93	34	2.70195086	0.803638546	-1.749382766	Down	1.03E-10	2.19E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012106 PE=4 SV=1
CCG002547.1	344	103	34	12.865938	3.455178518	-1.896724482	Down	6.19E-13	1.58E-12	AAEL000791-PA OS=Aedes aegypti GN=AAEL006791 PE=4 SV=1
CCG028532.1	2595	109	34	1.80489444	0.458027518	-1.97840828	Down	2.58E-14	8.98E-14	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001896 PE=4 SV=1
CCG014243.1	1227	110	34	3.85221745	0.968689006	-1.991583668	Down	1.51E-14	5.33E-14	AAEL003560-PA OS=Aedes aegypti GN=AAEL003560 PE=3 SV=1
CCG027763.1	1377	138	34	4.30633506	0.863167328	-2.318748412	Down	2.13E-21	8.40E-21	AGAP002322-PA OS=Anopheles gambiae GN=AgaP_AGAP002322 PE=4 SV=4
CCG000893.1	4767	141	34	1.2709739	0.249335307	-2.349775307	Down	3.65E-22	1.45E-21	AAEL003408-PA OS=Aedes aegypti GN=AAEL003408 PE=4 SV=1
CCG002665.1	1824	156	34	3.67504309	0.651634545	-2.495626174	Down	4.60E-26	1.89E-25	AAEL001203-PA OS=Aedes aegypti GN=AAEL001203 PE=4 SV=1
CCG015206.1	854	159	34	8.00021991	1.391781511	-2.52310691	Down	7.40E-27	3.05E-26	AAEL005485-PA OS=Aedes aegypti GN=AAEL005485 PE=4 SV=1
CCG007808.1	723	181	34	10.7572918	1.643957691	-2.710069842	Down	8.50E-33	3.63E-32	AGAP013096-PA OS=Anopheles gambiae GN=AgaP_AGAP013096 PE=4 SV=1
CCG001430.1	894	197	34	9.46872228	1.329509407	-2.832275774	Down	3.13E-37	1.37E-36	-
CCG024006.1	549	206	34	16.1234341	2.164993461	-2.896724482	Down	9.20E-40	4.10E-39	AAEL015151-PA OS=Aedes aegypti GN=AAEL015151 PE=4 SV=1
CCG012253.1	540	333	34	26.498003	2.201076686	-3.589602322	Down	2.26E-77	1.14E-76	AAEL003721-PA OS=Aedes aegypti GN=AAEL003721 PE=4 SV=1
CCG017704.1	307	0	35	0.001	3.985471293	11.96053462	Up	9.86E-10	2.03E-09	-

CCG022715.1	507	0	35	0.001	2.413293268	11.23678753	Up	9.86E-10	2.03E-09	AAEL014001-PA OS=Aedes aegypti GN=AAEL014001 PE=4 SV=1
CCG024788.1	606	0	35	0.001	2.019042388	10.97945548	Up	9.86E-10	2.03E-09	AAEL011908-PA OS=Aedes aegypti GN=AAEL011908 PE=3 SV=1
CCG006370.1	853	0	35	0.001	1.434395882	10.48622754	Up	9.86E-10	2.03E-09	-
CCG003594.1	1300	0	35	0.001	0.941184375	9.878333559	Up	9.86E-10	2.03E-09	Uncharacterized protein (Fragment) OS=Xenopus tropicalis GN=LOC100489570 PE=4 SV=1
CCG022958.1	633	1	35	0.06788268	1.932922096	4.831596221	Up	0.000587	0.0009151	AAEL000896-PA OS=Aedes aegypti GN=AAEL000896 PE=4 SV=1
CCG015907.1	567	1	35	0.07578436	2.157918319	4.831596221	Up	0.000587	0.00091504	AAEL007059-PA OS=Aedes aegypti GN=PGRPS5_b PE=4 SV=1
CCG026332.1	1124	2	35	0.0764586	1.08855844	3.831596221	Up	9.37E-05	0.00015325	AAEL007996-PA OS=Aedes aegypti GN=AAEL007996 PE=4 SV=1
CCG000475.1	479	3	35	0.26912151	2.554362603	3.24663372	Up	5.37E-05	8.96E-05	-
CCG005344.1	462	4	35	0.37203233	2.648354301	2.831596221	Up	3.37E-05	5.71E-05	AAEL010835-PA (Fragment) OS=Aedes aegypti GN=AAEL010835 PE=4 SV=1
CCG000095.1	1434	6	35	0.17978968	0.853235486	2.24663372	Up	6.05E-05	0.00010058	Predicted protein OS=Nematostella vectensis GN=v1g196134 PE=4 SV=1
CCG009549.1	1476	7	35	0.203786	0.828956428	2.024241299	Up	0.0001537	0.00024835	Retrotransposable element Tf2 155 kDa protein type, putative OS=Perkinsus marinus (strain ATCC 50983 / TXsc) GN=Pmar_PMAR017345 PE=4 SV=1
CCG006771.1	2412	7	35	0.12470487	0.507271844	2.024241299	Up	0.0001537	0.00024834	AAEL000522-PA OS=Aedes aegypti GN=AAEL000522 PE=4 SV=1
CCG008341.1	942	7	35	0.31930801	1.298874402	2.024241299	Up	0.0001537	0.00024837	AAEL002273-PA OS=Aedes aegypti GN=AAEL002273 PE=3 SV=1
CCG004770.1	779	8	35	0.44128097	1.570654284	1.831596221	Up	0.0003861	0.00061063	AGAP010462-PA (Fragment) OS=Anopheles gambiae GN=AGAP010462 PE=4 SV=3

CCG028264.1	1119	69	35	2.6496083	1.093422419	-1.276928236	Down	1.09E-05	1.91E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_02396 PE=4 SV=1
CCG024346.1	370	70	35	8.12940925	3.306864019	-1.297686796	Down	7.26E-06	1.29E-05	-
CCG001370.1	1056	78	35	3.17390085	1.158655007	-1.453805998	Down	2.37E-07	4.48E-07	AAEL004295-PA OS=Aedes aegypti GN=AAEL004295 PE=4 SV=1
CCG028362.1	915	106	35	4.97791461	1.337201844	-1.896324234	Down	2.88E-13	8.01E-13	AAEL012292-PA OS=Aedes aegypti GN=AAEL012292 PE=4 SV=1
CCG009800.1	849	129	35	6.52897028	1.441153931	-2.179631035	Down	1.02E-18	3.93E-18	AAEL005980-PA OS=Aedes aegypti GN=AAEL005980 PE=4 SV=1
CCG011866.1	1464	177	35	5.19511136	0.835751152	-2.636009329	Down	3.57E-31	1.51E-30	AAEL011364-PA OS=Aedes aegypti GN=AAEL011364 PE=4 SV=1
CCG017224.1	942	240	35	10.9477031	1.298874402	-3.075294375	Down	6.98E-49	3.24E-48	AAEL007852-PA OS=Aedes aegypti GN=AAEL007852 PE=4 SV=1
CCG008142.1	867	293	35	14.5214905	1.41123378	-3.363160634	Down	1.64E-64	8.01E-64	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG002588.1	327	322	35	42.312705	3.741711581	-3.499320657	Down	2.93E-73	1.47E-72	AAEL003247-PA OS=Aedes aegypti GN=AAEL003247 PE=4 SV=2
CCG026732.1	394	353	35	38.4982648	3.105430678	-3.631928153	Down	9.89E-83	5.05E-82	AAEL003843-PA OS=Aedes aegypti GN=AAEL003843 PE=4 SV=1
CCG000102.1	252	429	35	73.1508577	4.855316218	-3.913237617	Down	2.27E-106	1.21E-105	AAEL011157-PA OS=Aedes aegypti GN=AAEL011157 PE=4 SV=1
CCG004723.1	2079	576	35	11.9050347	0.588523178	-4.338328781	Down	3.07E-153	1.75E-152	AAEL015354-PA OS=Aedes aegypti GN=AAEL015354 PE=4 SV=1
CCG011427.1	576	0	36	0.001	2.184892298	11.09334645	Up	5.44E-10	1.13E-09	-
CCG026937.1	663	0	36	0.001	1.898186974	10.89040639	Up	5.44E-10	1.13E-09	AGAP006786-PA OS=Anopheles gambiae GN=AGAP006786 PE=4 SV=4
CCG014322.1	480	1	36	0.08952028	2.621870758	4.872238205	Up	0.000587	0.00091514	AAEL009770-PA OS=Aedes aegypti GN=AAEL009770 PE=2 SV=1
CCG024491.1	725	1	36	0.0592686	1.73585926	4.872238205	Up	0.000587	0.0009152	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG016271.1	1161	2	36	0.07402194	1.083977574	3.872238205	Up	9.36E-05	0.00015316	AAEL013852-PA OS=Aedes aegypti GN=AAEL013852 PE=4 SV=1

CCG004412.1	4492	2	36	0.01913167	0.280164284	3.872238205	Up	9.36E-05	0.00015318	BEL12_AG transposon polyprotein OS=Anopheles gambiae GN=BEL12-AG transposon PE=4 SV=1
CCG019174.1	399	2	36	0.21538714	3.154130235	3.872238205	Up	9.36E-05	0.00015315	AAEL002017-PA OS=Aedes aegypti GN=AAEL002617 PE=4 SV=1
CCG010622.1	1752	2	36	0.04905221	0.718320756	3.872238205	Up	9.36E-05	0.00015317	AAEL001986-PA OS=Aedes aegypti GN=AAEL001986 PE=3 SV=1
CCG009231.1	1629	3	36	0.07913395	0.772558603	3.287275705	Up	5.33E-05	8.90E-05	AAEL010759-PA OS=Aedes aegypti GN=AAEL010759 PE=4 SV=1
CCG022961.1	3178	4	36	0.054084	0.396003135	2.872238205	Up	3.21E-05	5.44E-05	RNase H and integrase-like protein (Fragment) OS=Bombyx mori GN=P-12 PE=4 SV=1
CCG006618.1	1563	4	36	0.10996733	0.805181039	2.872238205	Up	3.21E-05	5.44E-05	AAEL011888-PA OS=Aedes aegypti GN=AAEL011888 PE=3 SV=1
CCG021079.1	414	5	36	0.51895815	3.039850154	2.55031011	Up	2.59E-05	4.43E-05	-
CCG026215.1	462	6	36	0.5580485	2.724021567	2.287275705	Up	4.16E-05	7.01E-05	AAEL002253-PA OS=Aedes aegypti GN=AAEL002253 PE=4 SV=1
CCG020256.1	1353	6	36	0.19055315	0.930153706	2.287275705	Up	4.16E-05	7.01E-05	AAEL012594-PA (Fragment) OS=Aedes aegypti GN=AAEL012594 PE=4 SV=1
CCG024428.1	991	6	36	0.26015985	1.26992731	2.287275705	Up	4.16E-05	7.01E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_02250 PE=4 SV=1
CCG026607.1	448	7	36	0.6714021	2.809147241	2.064883283	Up	0.0001016	0.00016562	-
CCG013813.1	780	8	36	0.44071523	1.613458928	1.872238205	Up	0.0002575	0.0004118	AAEL014994-PA OS=Aedes aegypti GN=AAEL014994 PE=4 SV=1
CCG015251.1	868	8	36	0.39603442	1.449882447	1.872238205	Up	0.0002575	0.00041178	-
CCG000547.1	2926	8	36	0.1174839	0.430108668	1.872238205	Up	0.0002575	0.00041183	Phosphatidylinositol 3-kinase catalytic subunit type 3 OS=Trichinella spiralis GN=Tsp_00889 PE=4 SV=1

CCG007932.1	1220	9	36	0.31698985	1.031555708	1.702313204	Up	0.0006109	0.00095024	Putative uncharacterized protein GLEAN_15470 OS=Tribolium castaneum GN=GLEAN_15470 PE=4 SV=1
CCG003863.1	2301	60	36	1.12046244	0.54693523	-1.03465239	Down	0.0005359	0.00084035	AAEL005494-FA OS=Aedes aegypti GN=AAEL005494 PE=3 SV=1
CCG020444.1	1113	74	36	2.85692755	1.130725933	-1.33721516	Down	2.34E-06	4.23E-06	AAEL002671-FA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG011632.1	905	78	36	3.70346884	1.390605485	-1.413164014	Down	4.24E-07	7.93E-07	AAEL009929-FA OS=Aedes aegypti GN=AAEL009929 PE=4 SV=1
CCG012840.1	2193	81	36	1.58711742	0.573870481	-1.467611798	Down	1.13E-07	2.17E-07	AAEL000025-FA OS=Aedes aegypti GN=AAEL000025 PE=3 SV=1
CCG022500.1	2024	86	36	1.82578912	0.621787532	-1.554026549	Down	1.17E-08	2.33E-08	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010212 PE=4 SV=1
CCG026903.1	2217	87	36	1.68622775	0.56765808	-1.570705291	Down	7.34E-09	1.47E-08	Arylsulfatase B OS=Culex quinquefasciatus GN=CpipJ_CPIJ006774 PE=4 SV=1
CCG002841.1	1799	88	36	2.10191031	0.699554177	-1.587193413	Down	4.60E-09	9.29E-09	Ecdysis triggering hormone receptor subtype B OS=Aedes aegypti PE=2 SV=1
CCG021931.1	1317	90	36	2.93642833	0.95557932	-1.619614891	Down	1.79E-09	3.67E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002877 PE=4 SV=1
CCG015376.3	1525	91	36	2.56409564	0.825244566	-1.635556435	Down	1.11E-09	2.29E-09	AAEL009966-FA OS=Aedes aegypti GN=AAEL009966 PE=4 SV=1
CCG010637.3	2064	98	36	2.04022965	0.609737386	-1.742471639	Down	3.67E-11	7.96E-11	AAEL010044-FA OS=Aedes aegypti GN=AAEL010044 PE=4 SV=1
CCG020040.1	1830	99	36	2.3245922	0.687703805	-1.757118415	Down	2.23E-11	4.88E-11	AAEL004958-FA OS=Aedes aegypti GN=AAEL004958 PE=4 SV=1
CCG011188.1	1437	119	36	3.55838443	0.875781464	-2.022579558	Down	6.50E-16	2.38E-15	AAEL015017-FA OS=Aedes aegypti GN=AAEL015017 PE=4 SV=1
CCG023040.1	477	149	36	13.4224119	2.638360511	-2.346930315	Down	2.65E-23	1.06E-22	AAEL010529-FA OS=Aedes aegypti GN=AAEL010529 PE=4 SV=1

CCG010225.1	2678	149	36	2.39077314	0.469939494	-2.346930315	Down	2.65E-23	1.06E-22	AAEL012443-PA OS=Aedes aegypti GN=AAEL012443 PE=4 SV=1
CCG020358.1	1053	151	36	6.16185178	1.195154761	-2.366166534	Down	8.13E-24	3.27E-23	AAEL001427-PA OS=Aedes aegypti GN=AAEL001427 PE=4 SV=1
CCG024345.1	1188	157	36	5.67866021	1.05934172	-2.422382544	Down	2.29E-25	9.35E-25	AAEL005244-PA OS=Aedes aegypti GN=AAEL005244 PE=4 SV=1
CCG024545.1	2449	175	36	3.07052003	0.513882386	-2.578972907	Down	4.04E-30	1.70E-29	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010686 PE=4 SV=1
CCG028073.1	801	207	36	11.1045382	1.571158507	-2.821248752	Down	6.93E-39	3.08E-38	AAEL013483-PA OS=Aedes aegypti GN=AAEL013483 PE=4 SV=1
CCG008803.1	1842	218	36	5.08545176	0.68322365	-2.89594612	Down	5.63E-42	2.53E-41	AAEL014086-PA OS=Aedes aegypti GN=AAEL014086 PE=4 SV=1
CCG012223.1	1032	279	36	11.6168178	1.219474771	-3.251883107	Down	1.20E-59	5.78E-59	AAEL009295-PA OS=Aedes aegypti GN=AAEL009295 PE=4 SV=1
CCG028168.1	3699	358	36	4.15873614	0.340226538	-3.611577572	Down	1.74E-83	8.88E-83	AAEL004824-PA (Fragment) OS=Aedes aegypti GN=AAEL004824 PE=4 SV=1
CCG007757.1	1635	703	36	18.4756718	0.769723525	-4.585142674	Down	1.84E-193	1.09E-192	AAEL015128-PA OS=Aedes aegypti GN=AAEL015128 PE=4 SV=1
CCG007694.1	304	0	37	0.001	4.254790265	12.0548723	Up	3.00E-10	6.30E-10	Importin OS=Culex quinquefasciatus GN=CpipJ_CPIJ008197 PE=4 SV=1
CCG009214.1	2646	0	37	0.001	0.488834558	8.93320247	Up	3.00E-10	6.30E-10	AAEL009308-PA OS=Aedes aegypti GN=AAEL009308 PE=4 SV=1
CCG019605.1	1542	1	37	0.02786624	0.838817277	4.911766569	Up	0.000587	0.00091525	AAEL011028-PA OS=Aedes aegypti GN=AAEL011028 PE=4 SV=1
CCG010129.1	609	2	37	0.14111571	2.123901873	3.911766569	Up	9.36E-05	0.00015313	AAEL003927-PA OS=Aedes aegypti GN=AAEL003927 PE=4 SV=1
CCG021352.1	657	4	37	0.26161178	1.96873096	2.911766569	Up	3.11E-05	5.27E-05	AAEL000890-PA OS=Aedes aegypti GN=AAEL000890 PE=4 SV=1
CCG027996.1	1275	4	37	0.13480701	1.014475483	2.911766569	Up	3.11E-05	5.27E-05	AAEL000683-PA OS=Aedes aegypti GN=AAEL000683 PE=4 SV=1

CCG027407.4	3302	4	37	0.05205298	0.391719031	2.911766569	Up	3.11E-05	5.27E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC001426 PE=4 SV=1
CCG008206.1	869	6	37	0.29668401	1.488442164	2.326804069	Up	2.95E-05	5.03E-05	-
CCG017002.1	1632	6	37	0.15797697	0.792558971	2.326804069	Up	2.95E-05	5.03E-05	-
CCG017561.1	278	7	37	1.08197173	4.65272029	2.104411647	Up	6.74E-05	0.00011187	-
CCG012024.1	1110	8	37	0.30969178	1.165275892	1.911766569	Up	0.0001713	0.000276	-
CCG005680.1	879	9	37	0.43996315	1.471508806	1.741841568	Up	0.0004132	0.0006527	Uncharacterized protein OS=Anopheles darlingi GN=AND_05546 PE=4 SV=1
CCG016117.1	775	9	37	0.49900337	1.668975794	1.741841568	Up	0.0004132	0.00065274	-
CCG022082.2	1479	65	37	1.88846028	0.87454783	-1.110601244	Down	0.0001342	0.00021768	AGAP005610-PA OS=Anopheles gambiae GN=AGAP005610 PE=4 SV=4
CCG002532.1	399	66	37	7.10777565	3.241744964	-1.13262755	Down	9.24E-05	0.00015231	AAEL000536-PA OS=Aedes aegypti GN=AAEL000536 PE=4 SV=1
CCG007798.1	1425	66	37	1.99017718	0.90768859	-1.13262755	Down	9.24E-05	0.0001523	AAEL005123-PA (Fragment) OS=Aedes aegypti GN=AAEL005123 PE=4 SV=2
CCG011730.1	1782	78	37	1.88083013	0.725845253	-1.373635649	Down	7.44E-07	1.38E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006773 PE=3 SV=1
CCG023203.1	639	89	37	5.98483002	2.02418817	-1.563966862	Down	5.51E-09	1.11E-08	Uncharacterized protein OS=Anopheles darlingi GN=AND_09440 PE=4 SV=1
CCG015586.1	1146	96	37	3.59955892	1.128670367	-1.673195931	Down	1.99E-10	4.21E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014452 PE=4 SV=1
CCG010510.1	912	96	37	4.52312996	1.418263422	-1.673195931	Down	1.99E-10	4.21E-10	AAEL001496-PA OS=Aedes aegypti GN=AAEL001496 PE=4 SV=1
CCG027757.1	8649	104	37	0.51669007	0.149549802	-1.788673149	Down	3.81E-12	8.83E-12	AAEL000700-PA OS=Aedes aegypti GN=AAEL000700 PE=3 SV=1
CCG019067.1	1405	106	37	3.24184475	0.920609424	-1.816153885	Down	1.38E-12	3.35E-12	-
CCG024799.1	522	127	37	10.4543224	2.477885518	-2.076918117	Down	1.98E-17	7.52E-17	-

CCG017001.1	3871	140	37	1.55405912	0.334140078	-2.217516447	Down	1.34E-20	5.26E-20	Putative uncharacterized protein OS=Caenorhabditis remanei GN=CRE_17270 PE=4 SV=1
CCG028383.1	900	144	37	6.87515754	1.437173601	-2.258158432	Down	1.35E-21	5.34E-21	AAEL008092-PA OS=Aedes aegypti GN=AAEL008092 PE=4 SV=1
CCG013115.1	931	144	37	6.64623178	1.38931927	-2.258158432	Down	1.35E-21	5.34E-21	-
CCG002523.1	426	156	37	15.7353958	3.036282255	-2.373635649	Down	1.21E-24	4.91E-24	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002862 PE=4 SV=1
CCG019479.1	813	165	37	8.72079485	1.590967086	-2.454555645	Down	5.61E-27	2.32E-26	AAEL000571-PA OS=Aedes aegypti GN=AAEL000571 PE=4 SV=1
CCG005767.1	521	166	37	13.6909327	2.482641537	-2.463272862	Down	3.07E-27	1.27E-26	-
CCG002685.1	1794	182	37	4.35924844	0.720990101	-2.596028071	Down	1.73E-31	7.37E-31	AAEL000127-PA OS=Aedes aegypti GN=AAEL000127 PE=4 SV=1
CCG023925.1	813	194	37	10.2535406	1.590967086	-2.688146273	Down	9.68E-35	4.19E-34	AAEL012640-PA OS=Aedes aegypti GN=AAEL012646 PE=4 SV=1
CCG006642.1	4131	195	37	2.02834622	0.313109717	-2.695563744	Down	5.16E-35	2.24E-34	AAEL007374-PA OS=Aedes aegypti GN=AAEL007374 PE=4 SV=1
CCG019243.1	825	231	37	12.0315257	1.567825746	-2.939982472	Down	4.58E-45	2.09E-44	AAEL001396-PA OS=Aedes aegypti GN=AAEL001396 PE=4 SV=1
CCG001629.1	489	266	37	23.3741297	2.645104786	-3.143515866	Down	3.70E-55	1.76E-54	AAEL008458-PA OS=Aedes aegypti GN=AAEL008458 PE=4 SV=1
CCG007373.1	1812	286	37	6.78219873	0.713827947	-3.248104767	Down	4.89E-61	2.37E-60	AAEL007968-PA OS=Aedes aegypti GN=AAEL007968 PE=4 SV=1
CCG023974.1	1275	344	37	11.5934029	1.014475483	-3.514498185	Down	1.91E-78	9.64E-78	AAEL006994-PA OS=Aedes aegypti GN=AAEL006994 PE=4 SV=1
CCG000019.1	1062	404	37	16.3463021	1.217943729	-3.746444913	Down	6.91E-97	3.64E-96	AAEL004133-PA OS=Aedes aegypti GN=AAEL004133 PE=4 SV=1
CCG027841.1	1467	585	37	17.1351702	0.881701595	-4.280526245	Down	3.39E-154	1.93E-153	AAEL001766-PA OS=Aedes aegypti GN=AAEL001766 PE=4 SV=1
CCG008209.1	480	0	38	0.001	2.767530244	11.43438337	Up	1.65E-10	3.51E-10	AAEL006670-PA OS=Aedes aegypti GN=AAEL006670 PE=4 SV=1

CCG028396.1	629	0	38	0.001	2.111946768	11.04435776	Up	1.65E-10	3.51E-10	ATP-binding cassette sub-family G member 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ012372 PE=3 SV=1
CCG023853.1	1434	0	38	0.001	0.926369956	9.855444655	Up	1.65E-10	3.51E-10	AAEL006556-PA OS=Aedes aegypti GN=AAEL008538 PE=4 SV=1
CCG026407.1	1280	1	38	0.03357011	1.037823842	4.950240717	Up	0.000587	0.00091536	-
CCG025537.1	596	2	38	0.14419374	2.228883418	3.950240717	Up	9.36E-05	0.00015311	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG003231.1	438	2	38	0.19620883	3.032909857	3.950240717	Up	9.36E-05	0.00015313	Reverse transcriptase OS=Aedes aegypti GN=pol PE=4 SV=1
CCG015576.1	1591	2	38	0.05401601	0.834955699	3.950240717	Up	9.36E-05	0.00015312	-
CCG000531.1	797	3	38	0.16174304	1.666768529	3.365278217	Up	5.30E-05	8.85E-05	-
CCG006648.1	444	4	38	0.38711473	2.991924589	2.950240717	Up	3.05E-05	5.17E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_09855 PE=4 SV=1
CCG019103.1	588	4	38	0.29231112	2.259208363	2.950240717	Up	3.05E-05	5.17E-05	AAEL015271-PA OS=Aedes aegypti GN=AAEL015271 PE=3 SV=1
CCG000496.4	1740	7	38	0.17286675	0.763456619	2.142885795	Up	4.51E-05	7.58E-05	AAEL001522-PA OS=Aedes aegypti GN=AAEL001522 PE=4 SV=1
CCG015243.1	519	9	38	0.74513991	2.559565544	1.780315716	Up	0.0002783	0.00044376	AAEL013338-PA OS=Aedes aegypti GN=AAEL013338 PE=3 SV=1
CCG009948.1	444	10	38	0.96778682	2.991924589	1.628312622	Up	0.0006347	0.0009862	Odorant-binding protein 12 OS=Culex quinquefasciatus PE=2 SV=1
CCG015137.1	1922	10	38	0.22356782	0.6911626	1.628312622	Up	0.0006347	0.00098626	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG008601.1	710	65	38	3.93384894	1.871006362	-1.072127096	Down	0.0002067	0.00033199	-
CCG023576.1	1325	76	38	2.46467912	1.002576994	-1.297686796	Down	2.94E-06	5.28E-06	AAEL005787-PA OS=Aedes aegypti GN=AAEL005787 PE=3 SV=1
CCG000249.1	261	78	38	12.8415299	5.089710794	-1.335161502	Down	1.28E-06	2.35E-06	Putative uncharacterized protein GLEAN_07454 OS=Tribolium castaneum GN=GLEAN_07454 PE=4 SV=1

CCG005706.1	1203	79	38	2.8217864	1.104251469	-1.353540031	Down	8.43E-07	1.56E-06	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG013865.1	6722	80	38	0.51139226	0.197621916	-1.371687378	Down	5.52E-07	1.03E-06	Pol-like protein OS=Aedes aegypti PE=4 SV=1
CCG004293.1	723	90	38	5.34892962	1.837364478	-1.541612379	Down	6.56E-09	1.32E-08	AAEL005463-FA OS=Aedes aegypti GN=AAEL005485 PE=4 SV=1
CCG009907.1	1341	97	38	3.10817618	0.990614853	-1.649672125	Down	2.45E-10	5.16E-10	AAEL009043-FA OS=Aedes aegypti GN=AAEL009045 PE=4 SV=1
CCG000994.1	1317	120	38	3.91523778	1.008667059	-1.956649878	Down	2.09E-15	7.56E-15	AGAP004235-PC OS=Anopheles gambiae GN=AgaP_AGAP004235 PE=3 SV=1
CCG009764.1	2013	137	38	2.9244181	0.659917793	-2.147791366	Down	1.90E-19	7.38E-19	AAEL002622-FA OS=Aedes aegypti GN=AAEL002622 PE=4 SV=1
CCG012531.1	1569	139	38	3.8067515	0.846663172	-2.168700355	Down	6.17E-20	2.41E-19	AAEL002814-FA OS=Aedes aegypti GN=AAEL002814 PE=4 SV=1
CCG011195.1	2066	164	38	3.41095667	0.642988634	-2.407311287	Down	3.03E-26	1.25E-25	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017464 PE=4 SV=1
CCG005501.1	2673	189	38	3.03826406	0.496975128	-2.612001707	Down	7.41E-33	3.17E-32	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC001808 PE=4 SV=1
CCG016900.1	1527	234	38	6.58475304	0.869950568	-2.920124002	Down	2.55E-45	1.16E-44	AAEL007607-FA OS=Aedes aegypti GN=CYP4D38 PE=3 SV=1
CCG012678.1	1731	314	38	7.79462546	0.767426064	-3.344380032	Down	1.14E-68	5.62E-68	AAEL007253-FA OS=Aedes aegypti GN=AAEL007253 PE=4 SV=1
CCG000126.1	6780	351	38	2.22453936	0.195931345	-3.505086503	Down	7.99E-80	4.06E-79	AAEL006527-FA OS=Aedes aegypti GN=AAEL006527 PE=4 SV=1
CCG010712.1	4248	506	38	5.11833468	0.312715282	-4.032752857	Down	4.86E-128	2.69E-127	DCAF_L2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006873 PE=4 SV=1
CCG002875.1	4322	1062	38	10.5585049	0.307361064	-5.102327334	Down	0	0	AAEL007226-FA OS=Aedes aegypti GN=AAEL007226 PE=4 SV=1
CCG021309.1	699	43768	38	2690.557	1.900449953	-10.46734812	Down	0	0	AAEL001319-FA OS=Aedes aegypti GN=AAEL001319 PE=4 SV=1

CCG020714.1	453	0	39	0.001	3.009652967	11.55538143	Up	9.12E-11	1.95E-10	AAEL002550-PA OS=Aedes aegypti GN=AAEL002550 PE=4 SV=1
CCG012118.1	1520	0	39	0.001	0.896955786	9.808893061	Up	9.12E-11	1.95E-10	Pol polyprotein (Fragment) OS=Anopheles gambiae PE=4 SV=1
CCG004888.2	1446	1	39	0.02971628	0.942858087	4.987715423	Up	0.000587	0.0009153	AAEL012230-PA OS=Aedes aegypti GN=AAEL012230 PE=4 SV=1
CCG023387.1	780	2	39	0.11017881	1.747913839	3.987715423	Up	9.36E-05	0.00015312	-
CCG006004.1	1422	3	39	0.09065345	0.958771304	3.402752922	Up	5.29E-05	8.84E-05	AAEL003714-PA OS=Aedes aegypti GN=AAEL003714 PE=4 SV=1
CCG006351.1	435	3	39	0.296343	3.134190331	3.402752922	Up	5.29E-05	8.83E-05	AAEL002875-PA OS=Aedes aegypti GN=AAEL002875 PE=4 SV=1
CCG000416.1	342	3	39	0.3769275	3.986470158	3.402752922	Up	5.29E-05	8.84E-05	AAEL003922-PA OS=Aedes aegypti GN=AAEL003922 PE=4 SV=1
CCG008499.1	731	3	39	0.17634638	1.865079062	3.402752922	Up	5.29E-05	8.84E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004469 PE=4 SV=1
CCG021600.1	855	4	39	0.201028	1.594588063	2.987715423	Up	3.01E-05	5.11E-05	-
CCG005294.1	2130	4	39	0.08069434	0.640081124	2.987715423	Up	3.01E-05	5.11E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014312 PE=4 SV=1
CCG018487.1	1150	4	39	0.14945995	1.18554156	2.987715423	Up	3.01E-05	5.11E-05	Serine/threonine-protein kinase rio2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009507 PE=4 SV=1
CCG001964.1	573	5	39	0.37495405	2.379359152	2.665787328	Up	1.83E-05	3.15E-05	AAEL0008915-PA OS=Aedes aegypti GN=AAEL0008915 PE=4 SV=1
CCG005798.1	1027	6	39	0.25104032	1.327529498	2.402752922	Up	1.68E-05	2.90E-05	GK10815 OS=Drosophila willistoni GN=GK10815 PE=4 SV=1
CCG010207.1	1587	6	39	0.16245646	0.859088087	2.402752922	Up	1.68E-05	2.90E-05	AAEL001730-PA OS=Aedes aegypti GN=AAEL001730 PE=4 SV=1
CCG027151.1	2079	6	39	0.12401078	0.65578297	2.402752922	Up	1.68E-05	2.90E-05	AAEL013981-PA OS=Aedes aegypti GN=AAEL013981 PE=4 SV=1
CCG015654.1	1389	7	39	0.21655014	0.981549888	2.180360501	Up	3.06E-05	5.19E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017585 PE=4 SV=1

CCG024956.1	684	7	39	0.43974875	1.993235079	2.180360501	Up	3.06E-05	5.19E-05	-
CCG003968.1	831	8	39	0.41366772	1.640641148	1.987715423	Up	7.55E-05	0.0001249	GD15954 OS=Drosophila simulans GN=GD15954 PE=4 SV=1
CCG024376.1	1155	8	39	0.29762587	1.180409346	1.987715423	Up	7.55E-05	0.00012492	Uncharacterized protein OS=Anopheles darlingi GN=AND_22299 PE=4 SV=1
CCG007550.1	337	8	39	1.02005305	4.0456166	1.987715423	Up	7.55E-05	0.00012491	AAEL010576-FJ OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG009173.1	1687	8	39	0.20376875	0.808164075	1.987715423	Up	7.55E-05	0.00012489	Putative uncharacterized protein OS=Caenorhabditis brenneri GN=CAEBREN_12673 PE=4 SV=1
CCG013189.1	1033	9	39	0.37437329	1.319818775	1.817790421	Up	0.0001868	0.00030071	-
CCG025808.1	454	10	39	0.94646993	3.003023776	1.665787328	Up	0.0004335	0.00068402	AAEL010576-FN OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG028085.1	626	10	39	0.68641749	2.177911812	1.665787328	Up	0.0004335	0.00068398	-
CCG021826.1	1005	73	39	3.1211847	1.356589845	-1.202109136	Down	1.61E-05	2.78E-05	AAEL015307-FA OS=Aedes aegypti GN=AAEL015307 PE=4 SV=1
CCG001962.1	1266	80	39	2.71530708	1.076913739	-1.334212672	Down	9.50E-07	1.75E-06	AAEL008186-FA OS=Aedes aegypti GN=AAEL008186 PE=4 SV=1
CCG003216.1	1434	82	39	2.45712569	0.950748113	-1.369836582	Down	4.09E-07	7.65E-07	AAEL005004-FA OS=Aedes aegypti GN=AAEL005004 PE=4 SV=1
CCG009151.1	813	85	39	4.49253068	1.676965306	-1.421675513	Down	1.12E-07	2.16E-07	AAEL005774-FA OS=Aedes aegypti GN=AAEL005774 PE=4 SV=1
CCG023158.1	993	111	39	4.80326339	1.37298368	-1.806700444	Down	4.99E-13	1.30E-12	AAEL013502-FA OS=Aedes aegypti GN=AAEL013502 PE=4 SV=1
CCG014544.1	2081	119	39	2.45718329	0.655152712	-1.907102341	Down	8.05E-15	2.89E-14	AAEL004862-FA OS=Aedes aegypti GN=AAEL004862 PE=4 SV=1
CCG017808.1	1041	144	39	5.94394023	1.309676075	-2.182209579	Down	9.45E-21	3.71E-20	-
CCG025932.1	1539	180	39	5.02569995	0.885882257	-2.504137674	Down	5.96E-30	2.51E-29	C-AMP specific 3',5'-cyclic phosphodiesterase OS=Culex quinquefasciatus GN=CpipJ_CPIJ003154 PE=3 SV=1
CCG015089.1	486	186	39	16.4452071	2.805293815	-2.551443388	Down	1.53E-31	6.48E-31	AAEL007493-FA OS=Aedes aegypti GN=AAEL007493 PE=4 SV=1

CCG006559.1	2103	193	39	3.94348967	0.648298999	-2.604741615	Down	2.03E-33	8.71E-33	AAEL000134-PA OS=Aedes aegypti GN=AAEL000134 PE=4 SV=1
CCG014923.1	462	244	39	22.6939724	2.951023364	-2.943021915	Down	1.46E-47	6.74E-47	AAEL001985-PA OS=Aedes aegypti GN=AAEL001985 PE=4 SV=1
CCG000890.1	219	359	39	70.4389713	6.225446548	-3.500124611	Down	1.66E-81	8.48E-81	AAEL003417-PA OS=Aedes aegypti GN=AAEL003417 PE=4 SV=1
CCG007814.1	648	492	39	32.6251689	2.103970361	-3.954799083	Down	9.85E-123	5.41E-122	AAEL009585-PA OS=Aedes aegypti GN=AAEL009585 PE=4 SV=1
CCG009557.1	1574	0	40	0.001	0.888393311	9.79505472	Up	5.03E-11	1.08E-10	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG009270.1	1695	2	40	0.05070175	0.824974083	4.024241299	Up	9.36E-05	0.00015312	AAEL003187-PA (Fragment) OS=Aedes aegypti GN=AAEL003187 PE=4 SV=1
CCG009045.1	516	4	40	0.33309872	2.709943936	3.024241299	Up	2.99E-05	5.07E-05	-
CCG022448.1	708	5	40	0.30345858	1.975043885	2.702313204	Up	1.74E-05	3.00E-05	AAEL000781-PA OS=Aedes aegypti GN=AAEL000781 PE=4 SV=1
CCG008463.1	1133	6	40	0.22755376	1.234184529	2.439278798	Up	1.37E-05	2.38E-05	AAEL014001-PA OS=Aedes aegypti GN=AAEL014001 PE=4 SV=1
CCG012654.1	1829	7	40	0.16445497	0.764533117	2.216886377	Up	2.12E-05	3.64E-05	AAEL005170-PA OS=Aedes aegypti GN=AAEL005170 PE=4 SV=1
CCG002442.1	594	68	40	4.9190942	2.354092712	-1.063221543	Down	0.0001633	0.00026344	AAEL011787-PA OS=Aedes aegypti GN=AAEL011787 PE=4 SV=1
CCG004470.1	2208	71	40	1.38172607	0.633302115	-1.125505821	Down	5.46E-05	9.11E-05	AAEL009062-PA OS=Aedes aegypti GN=AAEL009062 PE=4 SV=2
CCG004907.1	2307	71	40	1.32243223	0.606125302	-1.125505821	Down	5.46E-05	9.11E-05	Polyprotein OS=Antheraea mylitta PE=4 SV=1
CCG021943.1	408	73	40	7.68821232	3.427282036	-1.16558326	Down	2.57E-05	4.40E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_07352 PE=4 SV=1
CCG019476.1	774	73	40	4.05270107	1.806629291	-1.16558326	Down	2.57E-05	4.40E-05	AGAP002178-PA OS=Anopheles gambiae GN=AgaP_AGAP002178 PE=3 SV=5
CCG020813.1	717	74	40	4.43481222	1.95025254	-1.185212067	Down	1.75E-05	3.02E-05	-

CCG001626.1	2181	76	40	1.49734059	0.641142169	-1.223686215	Down	8.03E-06	1.42E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015620 PE=4 SV=1
CCG011834.1	1052	77	40	3.14512316	1.329212045	-1.242545242	Down	5.41E-06	9.60E-06	Hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG009265.1	3512	80	40	0.97880944	0.39815805	-1.297686796	Down	1.61E-06	2.94E-06	Pol polyprotein (Fragment) OS=Anopheles gambiae PE=4 SV=1
CCG010139.1	1884	88	40	2.0070789	0.742213944	-1.43519032	Down	5.40E-08	1.05E-07	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG025233.1	720	93	40	5.55025739	1.942126487	-1.514917512	Down	5.81E-09	1.17E-08	AAEL001235-PA OS=Aedes aegypti GN=AAEL001235 PE=4 SV=1
CCG015421.1	1080	106	40	4.21739988	1.294750992	-1.703679156	Down	1.25E-11	2.78E-11	AAEL001075-PA OS=Aedes aegypti GN=AAEL001075 PE=4 SV=1
CCG018764.1	855	110	40	5.52826995	1.635474937	-1.757118415	Down	1.74E-12	4.17E-12	AAEL012832-PA OS=Aedes aegypti GN=AAEL012832 PE=4 SV=1
CCG011358.1	1116	130	40	5.00543504	1.252984831	-1.998126514	Down	5.35E-17	2.02E-16	AAEL004213-PA OS=Aedes aegypti GN=AAEL004213 PE=4 SV=1
CCG007374.1	2222	135	40	2.61067244	0.629311913	-2.052574298	Down	3.55E-18	1.36E-17	AAEL000771-PA OS=Aedes aegypti GN=AAEL000771 PE=4 SV=1
CCG009075.1	2592	137	40	2.27116267	0.53947958	-2.073790784	Down	1.19E-18	4.56E-18	AAEL002754-PA OS=Aedes aegypti GN=AAEL002754 PE=4 SV=1
CCG024148.1	381	145	40	16.3533111	3.670160291	-2.155667791	Down	1.38E-20	5.43E-20	-
CCG010121.1	1983	147	40	3.18535098	0.70515939	-2.175431046	Down	4.49E-21	1.77E-20	AAEL012091-PA OS=Aedes aegypti GN=AAEL012091 PE=3 SV=1
CCG000099.1	357	147	40	17.6934201	3.916893756	-2.175431046	Down	4.49E-21	1.77E-20	-
CCG021729.1	1503	156	40	4.45993253	0.930359994	-2.26116092	Down	2.63E-23	1.05E-22	AAEL012065-PA OS=Aedes aegypti GN=AAEL012065 PE=4 SV=1
CCG019873.1	543	160	40	12.6614319	2.575195342	-2.297686796	Down	2.59E-24	1.05E-23	-
CCG013166.1	388	164	40	18.1624651	3.603946059	-2.333310706	Down	2.50E-25	1.02E-24	-
CCG018913.1	1347	166	40	5.29545356	1.038107699	-2.350798133	Down	7.73E-26	3.16E-25	AAEL013648-PA OS=Aedes aegypti GN=AAEL013648 PE=4 SV=1
CCG003109.1	930	176	40	8.13190677	1.503581797	-2.43519032	Down	2.03E-28	8.46E-28	AAEL002088-PA OS=Aedes aegypti GN=AAEL002088 PE=4 SV=1

CCG014710.1	3438	178	40	2.22472739	0.40672806	-2.451492132	Down	6.09E-29	2.56E-28	AAEL009258-PA OS=Aedes aegypti GN=AAEL009258 PE=4 SV=1
CCG019115.1	1980	256	40	5.55568286	0.706227814	-2.975758701	Down	2.23E-50	1.04E-49	AAEL002173-PA OS=Aedes aegypti GN=AAEL002173 PE=4 SV=1
CCG011272.1	1012	258	40	10.9547347	1.38175007	-2.986985957	Down	5.99E-51	2.80E-50	-
CCG015508.1	1166	258	40	9.50788296	1.199254778	-2.986985957	Down	5.99E-51	2.80E-50	Homeotic protein span-major OS=Culex quinquefasciatus GN=CpipJ_CPIJ015319 PE=4 SV=1
CCG015542.1	1704	292	40	7.36335828	0.820616826	-3.16558326	Down	8.52E-61	4.13E-60	AAEL005386-PA OS=Aedes aegypti GN=AAEL005386 PE=4 SV=1
CCG015945.1	573	1691	40	126.809461	2.440368361	-5.699419646	Down	0	0	AAEL008285-PA OS=Aedes aegypti GN=AAEL008285 PE=4 SV=1
CCG006476.1	704	0	41	0.001	2.035922369	10.99146684	Up	2.77E-11	6.04E-11	AGAP008251-PA OS=Anopheles gambiae GN=AGAP008251 PE=4 SV=3
CCG024174.1	1173	0	41	0.001	1.221900552	10.25491116	Up	2.77E-11	6.04E-11	AAEL009292-PA OS=Aedes aegypti GN=AAEL009292 PE=4 SV=1
CCG005660.1	591	2	41	0.14541365	2.425193482	4.059865208	Up	9.36E-05	0.00015312	AAEL013345-PA OS=Aedes aegypti GN=AAEL013345 PE=3 SV=1
CCG017625.1	435	3	41	0.296343	3.294918041	3.474902708	Up	5.28E-05	8.83E-05	AAEL002446-PA (Fragment) OS=Aedes aegypti GN=AAEL002446 PE=4 SV=1
CCG016386.2	435	3	41	0.296343	3.294918041	3.474902708	Up	5.28E-05	8.82E-05	AAEL005474-PA OS=Aedes aegypti GN=AAEL005474 PE=4 SV=1
CCG027118.1	1026	4	41	0.16752333	1.396968175	3.059865208	Up	2.97E-05	5.05E-05	Transcription initiation factor IIF large subunit OS=Aedes aegypti PE=2 SV=1
CCG028057.1	1746	4	41	0.09844155	0.820898825	3.059865208	Up	2.97E-05	5.05E-05	-
CCG001448.1	527	6	41	0.48921899	2.719714132	2.474902708	Up	1.17E-05	2.04E-05	-
CCG006936.2	830	7	41	0.36239535	1.726854636	2.252510286	Up	1.51E-05	2.62E-05	AAEL006672-PA (Fragment) OS=Aedes aegypti GN=AAEL006672 PE=4 SV=1
CCG007759.1	945	7	41	0.31829433	1.516708304	2.252510286	Up	1.51E-05	2.62E-05	AAEL010768-PA OS=Aedes aegypti GN=AAEL010768 PE=3 SV=1
CCG016204.1	462	7	41	0.65105659	3.102357895	2.252510286	Up	1.51E-05	2.62E-05	-

CCG019409.1	1569	8	41	0.21909361	0.913505002	2.059865208	Up	3.35E-05	5.67E-05	AGAP001625-PA OS=Anopheles gambiae GN=AgaP_AGAP001625 PE=4 SV=5
CCG003916.1	927	8	41	0.37082835	1.546158951	2.059865208	Up	3.35E-05	5.67E-05	AAEL011493-PA OS=Aedes aegypti GN=AAEL011493 PE=4 SV=1
CCG018359.1	1266	8	41	0.27153071	1.132140085	2.059865208	Up	3.35E-05	5.67E-05	Four1 gag/pol protein OS=Zea mays GN=Fourf gag/pol PE=4 SV=1
CCG008129.1	4449	9	41	0.08692461	0.322159889	1.889940207	Up	8.35E-05	0.0001377	AAEL004216-PA OS=Aedes aegypti GN=AAEL004216 PE=4 SV=1
CCG007965.1	1392	10	41	0.30869062	1.029661888	1.737937114	Up	0.0001998	0.00032121	-
CCG024133.1	1263	10	41	0.34021959	1.134829254	1.737937114	Up	0.0001998	0.00032123	AAEL006457-PA OS=Aedes aegypti GN=AAEL006457 PE=4 SV=1
CCG022678.1	1755	11	41	0.26932597	0.816689087	1.60043359	Up	0.0004474	0.00070523	AAEL013804-PA OS=Aedes aegypti GN=AAEL013804 PE=4 SV=1
CCG025727.1	441	11	41	1.07180744	3.250089224	1.60043359	Up	0.0004474	0.00070527	AAEL004609-PA OS=Aedes aegypti GN=AAEL004609 PE=4 SV=1
CCG011595.1	1002	86	41	3.68802113	1.430428491	-1.366399546	Down	2.25E-07	4.26E-07	AAEL002116-PA OS=Aedes aegypti GN=AAEL002116 PE=4 SV=1
CCG009405.1	1176	106	41	3.87312234	1.218783459	-1.668055246	Down	2.52E-11	5.51E-11	AAEL014658-PA OS=Aedes aegypti GN=AAEL014658 PE=3 SV=1
CCG019229.1	1469	108	41	3.15910915	0.975690502	-1.695022294	Down	9.57E-12	2.13E-11	AAEL007811-PA OS=Aedes aegypti GN=AAEL007811 PE=2 SV=1
CCG021137.1	1356	110	41	3.48574543	1.056998044	-1.721494505	Down	3.59E-12	8.35E-12	Protease (Reverse transcriptase) (Rnase h) (Integrase) (Fragment) OS=Drosophila melanogaster GN=pol PE=4 SV=1
CCG003991.1	993	113	41	4.88980867	1.443393099	-1.760313754	Down	8.12E-13	2.03E-12	AAEL011454-PA OS=Aedes aegypti GN=AAEL011454 PE=4 SV=1
CCG000212.1	762	126	41	7.10523171	1.880957149	-1.917414715	Down	1.04E-15	3.77E-15	AGAP001753-PA OS=Anopheles gambiae GN=AGAP001753 PE=3 SV=4
CCG015351.1	888	128	41	6.19383562	1.614064581	-1.940134792	Down	3.61E-16	1.35E-15	AAEL009876-PA OS=Aedes aegypti GN=AAEL009876 PE=3 SV=1
CCG002014.1	1107	133	41	5.16257877	1.294750992	-1.995417227	Down	2.51E-17	9.51E-17	AAEL001262-PA OS=Aedes aegypti GN=AAEL001262 PE=4 SV=1

CCG010416.1	1140	140	41	5.27698495	1.257271358	-2.069417809	Down	5.57E-19	2.15E-18	AAEL004935-PA OS=Aedes aegypti GN=AAEL004935 PE=4 SV=1
CCG025428.1	2937	143	41	2.09215936	0.488011354	-2.100006128	Down	1.07E-19	4.14E-19	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002887 PE=4 SV=1
CCG021356.1	2043	156	41	3.28109574	0.70156111	-2.22553701	Down	7.05E-23	2.82E-22	AAEL002880-PA OS=Aedes aegypti GN=AAEL002880 PE=3 SV=1
CCG023110.1	528	182	41	14.8115373	2.714563158	-2.447929432	Down	1.65E-29	6.94E-29	AAEL005899-PA OS=Aedes aegypti GN=AAEL005899 PE=4 SV=1
CCG014880.1	2070	195	41	4.04787355	0.692410313	-2.547465105	Down	6.15E-33	2.63E-32	AAEL011764-PA OS=Aedes aegypti GN=PPO10 PE=4 SV=1
CCG008103.1	2991	208	41	2.98819953	0.479200718	-2.64057451	Down	1.98E-36	8.66E-36	AAEL000778-PA OS=Aedes aegypti GN=AAEL000778 PE=4 SV=1
CCG001154.1	1284	442	41	14.7917622	1.116268962	-3.728037351	Down	2.02E-105	1.08E-104	AAEL014445-PA OS=Aedes aegypti GN=AAEL014445 PE=4 SV=1
CCG015422.1	312	7820	41	1076.99784	4.593876114	-7.873087684	Down	0	0	AAEL001082-PA OS=Aedes aegypti GN=AAEL001082 PE=4 SV=1
CCG002499.1	561	8603	41	658.945859	2.554882973	-8.010758914	Down	0	0	AAEL001683-PA OS=Aedes aegypti GN=AAEL001683 PE=4 SV=1
CCG006563.1	375	0	42	0.001	3.915326998	11.93491709	Up	1.53E-11	3.37E-11	AAEL000138-PA OS=Aedes aegypti GN=AAEL000138 PE=4 SV=1
CCG016735.1	381	0	42	0.001	3.853668306	11.91201669	Up	1.53E-11	3.37E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004470 PE=4 SV=1
CCG001867.2	789	0	42	0.001	1.860896862	10.86178238	Up	1.53E-11	3.37E-11	AAEL008206-PA OS=Aedes aegypti GN=AAEL008206 PE=3 SV=1
CCG003928.1	1276	0	42	0.001	1.150664282	10.16825126	Up	1.53E-11	3.37E-11	AAEL012229-PA OS=Aedes aegypti GN=AAEL012229 PE=4 SV=1
CCG002600.1	1395	0	42	0.001	1.052507258	10.03961447	Up	1.53E-11	3.37E-11	Beta-galactosidase OS=Aedes aegypti GN=AAEL004575 PE=3 SV=1
CCG010902.1	4749	0	42	0.001	0.309169851	8.272255832	Up	1.53E-11	3.37E-11	AAEL008384-PA OS=Aedes aegypti GN=AAEL008384 PE=3 SV=1
CCG020265.1	336	1	42	0.12788611	4.369784597	5.094630627	Up	0.000587	0.00091848	-

CCG026612.1	2073	2	42	0.04145657	0.708271888	4.094630627	Up	9.36E-05	0.00015313	Polyribonucleotide nucleotidyltransferase OS=Wolbachia pipientis wAlbB GN=ppn PE=3 SV=1
CCG026267.1	516	3	42	0.24982404	2.845441133	3.509668126	Up	5.28E-05	8.83E-05	AAEL010115-FA OS=Aedes aegypti GN=AAEL010115 PE=4 SV=1
CCG011317.1	501	3	42	0.2573038	2.930633981	3.509668126	Up	5.28E-05	8.83E-05	AAEL002679-FA OS=Aedes aegypti GN=AAEL002679 PE=4 SV=1
CCG000654.1	1566	6	42	0.164635	0.937578304	2.509668126	Up	1.05E-05	1.83E-05	AAEL013489-FA OS=Aedes aegypti GN=AAEL013489 PE=4 SV=1
CCG005619.1	555	7	42	0.54196062	2.645491215	2.287275705	Up	1.12E-05	1.95E-05	AAEL010576-FA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG006743.1	666	8	42	0.51615297	2.204576013	2.094630627	Up	2.25E-05	3.87E-05	AAEL003678-FA OS=Aedes aegypti GN=AAEL003678 PE=3 SV=1
CCG026528.1	901	8	42	0.38152928	1.62957561	2.094630627	Up	2.25E-05	3.87E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016207 PE=4 SV=1
CCG009721.2	1605	8	42	0.21417936	0.914796028	2.094630627	Up	2.25E-05	3.87E-05	AAEL002349-FA OS=Aedes aegypti GN=AAEL002349 PE=4 SV=1
CCG012292.1	804	10	42	0.53444944	1.826178637	1.772702532	Up	0.0001349	0.00021874	-
CCG006637.1	2413	10	42	0.17807598	0.608473943	1.772702532	Up	0.0001349	0.00021873	AAEL014411-FA OS=Aedes aegypti GN=CYP304B3 PE=3 SV=1
CCG008067.1	496	11	42	0.95295782	2.960176662	1.635199008	Up	0.0003071	0.00048823	Putative uncharacterized protein OS=Aedes albopictus PE=2 SV=1
CCG017265.1	351	69	42	8.44704185	4.183041665	-1.01389383	Down	0.00026	0.00041577	AAEL006854-FA OS=Aedes aegypti GN=AAEL006854 PE=4 SV=1
CCG000984.1	3125	70	42	0.96252206	0.46983924	-1.03465239	Down	0.0001836	0.00029561	5-hydroxytryptamine 7 receptor OS=Aedes aegypti PE=2 SV=1
CCG004113.1	663	72	42	4.66639652	2.21455147	-1.075294375	Down	9.02E-05	0.00014863	-
CCG026599.1	1023	78	42	3.27628475	1.43523717	-1.190771592	Down	9.60E-06	1.68E-05	AAEL015331-FA OS=Aedes aegypti GN=AAEL015331 PE=4 SV=1
CCG015745.1	2579	81	42	1.3495729	0.569308889	-1.245219376	Down	2.96E-06	5.33E-06	AAEL005669-FA OS=Aedes aegypti GN=AAEL005669 PE=4 SV=1
CCG028638.1	1571	89	42	2.43431342	0.934594287	-1.381102804	Down	1.09E-07	2.09E-07	GK10815 OS=Drosophila willistoni GN=GK10815 PE=4 SV=1

CCG022383.1	543	103	42	8.1507968	2.703955109	-1.591869901	Down	2.06E-10	4.35E-10	AAEL011859-PA OS=Aedes aegypti GN=AAEL011859 PE=4 SV=1
CCG012819.1	2928	112	42	1.64365105	0.501450691	-1.712724295	Down	2.75E-12	6.46E-12	AAEL003305-PA OS=Aedes aegypti GN=AAEL003305 PE=4 SV=1
CCG014993.1	702	132	42	8.07977916	2.091520833	-1.949763493	Down	9.93E-17	3.74E-16	AAEL002447-PA OS=Aedes aegypti GN=AAEL002447 PE=4 SV=1
CCG000299.1	1752	140	42	3.43365459	0.838040882	-2.03465239	Down	1.35E-18	5.19E-18	AAEL009856-PA OS=Aedes aegypti GN=AAEL009856 PE=4 SV=1
CCG015044.1	1713	141	42	3.53691336	0.857120621	-2.044920726	Down	7.82E-19	3.01E-18	AAEL005345-PA OS=Aedes aegypti GN=AAEL005345 PE=4 SV=1
CCG002700.1	1089	150	42	5.91869623	1.348253099	-2.134188064	Down	5.43E-21	2.14E-20	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004376 PE=4 SV=1
CCG016334.1	1248	187	42	6.43857402	1.176480468	-2.452263833	Down	2.45E-30	1.03E-29	AAEL003093-PA OS=Aedes aegypti GN=AAEL003093 PE=4 SV=1
CCG015680.1	1943	188	42	4.15764802	0.755660126	-2.459958225	Down	1.34E-30	5.67E-30	AAEL008483-PA OS=Aedes aegypti GN=AAEL008483 PE=4 SV=1
CCG000972.1	1626	224	42	5.91956984	0.902981319	-2.712724295	Down	2.94E-40	1.31E-39	AAEL011699-PA OS=Aedes aegypti GN=AAEL011699 PE=4 SV=1
CCG024739.1	2370	239	42	4.33323484	0.619513766	-2.806236181	Down	2.09E-44	9.53E-44	AAEL008831-PA OS=Aedes aegypti GN=AAEL008831 PE=4 SV=1
CCG026524.1	1128	350	42	13.3328077	1.301637965	-3.356580485	Down	1.31E-76	6.62E-76	AAEL009359-PA OS=Aedes aegypti GN=AAEL009359 PE=4 SV=1
CCG025866.1	498	373	42	32.1841587	2.948288402	-3.448401194	Down	1.59E-83	8.15E-83	AAEL008069-PA OS=Aedes aegypti GN=AAEL008069 PE=4 SV=1
CCG007865.1	888	698	42	33.7757599	1.65343201	-4.3524526	Down	9.60E-186	5.67E-185	AAEL000762-PA OS=Aedes aegypti GN=AAEL000762 PE=4 SV=1
CCG012033.1	315	0	43	0.001	4.772082226	12.22040319	Up	8.43E-12	1.88E-11	AAEL013514-PA OS=Aedes aegypti GN=AAEL013514 PE=4 SV=1
CCG010138.1	375	0	43	0.001	4.00854907	11.96886442	Up	8.43E-12	1.88E-11	Chromosome undetermined SCAF14565, whole genome shotgun sequence OS=Tetraodon nigroviridis GN=GSTENG00016696001 PE=3 SV=1

CCG011485.1	787	0	43	0.001	1.910045618	10.89939138	Up	8.43E-12	1.88E-11	-
CCG005993.1	810	0	43	0.001	1.855809755	10.85783311	Up	8.43E-12	1.88E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017971 PE=4 SV=1
CCG004266.1	993	0	43	0.001	1.513802519	10.5639613	Up	8.43E-12	1.88E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005433 PE=4 SV=1
CCG002818.1	709	1	43	0.06060611	2.120177576	5.128577959	Up	0.000587	0.00091575	-
CCG019318.1	471	1	43	0.09123086	3.19151996	5.128577959	Up	0.000587	0.0009166	AAEL000670-PA OS=Aedes aegypti GN=AAEL006670 PE=4 SV=1
CCG011361.1	1464	1	43	0.02935091	1.026779987	5.128577959	Up	0.000587	0.00091825	AAEL004200-PA OS=Aedes aegypti GN=AAEL004206 PE=4 SV=1
CCG024031.1	633	2	43	0.13576535	2.374732861	4.128577959	Up	9.36E-05	0.00015313	AAEL004662-PA OS=Aedes aegypti GN=AAEL004662 PE=4 SV=1
CCG020209.1	2046	3	43	0.06300548	0.734704742	3.543615458	Up	5.28E-05	8.83E-05	AAEL002537-PA OS=Aedes aegypti GN=AAEL002537 PE=4 SV=1
CCG012437.1	639	3	43	0.20173584	2.3524349	3.543615458	Up	5.28E-05	8.83E-05	AAEL007809-PA OS=Aedes aegypti GN=AAEL007809 PE=4 SV=1
CCG021924.1	519	5	43	0.41396661	2.896350484	2.806649864	Up	1.63E-05	2.81E-05	AAEL000267-PB OS=Aedes aegypti GN=AAEL000267 PE=4 SV=1
CCG006735.1	973	5	43	0.22081056	1.544918706	2.806649864	Up	1.63E-05	2.81E-05	-
CCG022311.1	1137	6	43	0.22675322	1.322080828	2.543615458	Up	9.67E-06	1.69E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004227 PE=4 SV=1
CCG012211.1	1233	7	43	0.24394821	1.219145094	2.321223036	Up	8.69E-06	1.52E-05	AAEL000285-PA OS=Aedes aegypti GN=AAEL000285 PE=4 SV=1
CCG016592.1	660	8	43	0.52084527	2.277584699	2.128577959	Up	1.54E-05	2.66E-05	AAEL000414-PA OS=Aedes aegypti GN=AAEL000414 PE=3 SV=1
CCG010253.1	1218	8	43	0.28223143	1.234159196	2.128577959	Up	1.54E-05	2.66E-05	AAEL000009-PA OS=Aedes aegypti GN=AAEL000009 PE=4 SV=1
CCG014439.1	977	9	43	0.39583174	1.538593553	1.958652957	Up	3.71E-05	6.26E-05	-
CCG012173.1	1880	11	43	0.25141866	0.799577607	1.66914634	Up	0.0002099	0.00033707	AAEL004635-PA OS=Aedes aegypti GN=AAEL004635 PE=4 SV=1

CCG012234.1	1950	11	43	0.24239337	0.770874821	1.66914634	Up	0.0002099	0.00033702	AAEL010512-PA OS=Aedes aegypti GN=AAEL010512 PE=4 SV=1
CCG028424.1	2501	11	43	0.18899124	0.601041944	1.66914634	Up	0.0002099	0.000337	BEL12_AG transposon polyprotein OS=Anopheles gambiae GN=BEL12-AG transposon PE=4 SV=1
CCG025339.1	2357	11	43	0.20053758	0.637762368	1.66914634	Up	0.0002099	0.00033705	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004811 PE=4 SV=1
CCG004864.1	1650	12	43	0.31250716	0.91103388	1.543615458	Up	0.0004553	0.0007173	AAEL001996-PA OS=Aedes aegypti GN=AAEL001996 PE=4 SV=1
CCG003790.1	878	12	43	0.58728567	1.712079614	1.543615458	Up	0.0004553	0.00071735	AAEL0008790-PA OS=Aedes aegypti GN=AAEL008790 PE=4 SV=1
CCG012441.1	1149	77	43	2.87960798	1.308273195	-1.138208582	Down	2.23E-05	3.82E-05	AAEL011987-PA OS=Aedes aegypti GN=AAEL011987 PE=4 SV=1
CCG024689.1	1347	78	43	2.48822517	1.115965777	-1.15682426	Down	1.53E-05	2.65E-05	AAEL000250-PA OS=Aedes aegypti GN=AAEL000250 PE=4 SV=1
CCG006568.1	324	79	43	10.4771884	4.639524386	-1.17520279	Down	1.04E-05	1.83E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014977 PE=4 SV=1
CCG021360.1	1677	90	43	2.30606805	0.896366071	-1.363275138	Down	1.24E-07	2.37E-07	AAEL009616-PA OS=Aedes aegypti GN=AAEL009616 PE=4 SV=1
CCG004831.1	393	130	43	14.2139071	3.824951403	-1.893789855	Down	6.43E-16	2.35E-15	AAEL010108-PA OS=Aedes aegypti GN=AAEL010108 PE=4 SV=1
CCG007841.1	420	139	43	14.220936	3.57906167	-1.990363114	Down	5.49E-18	2.10E-17	AAEL002811-PA OS=Aedes aegypti GN=AAEL002811 PE=4 SV=1
CCG004846.1	903	151	43	7.1854152	1.664679846	-2.109826781	Down	7.81E-21	3.07E-20	AAEL0006477-PA OS=Aedes aegypti GN=AAEL006477 PE=4 SV=1
CCG018838.1	1215	157	43	5.55246776	1.237206503	-2.16604279	Down	2.73E-22	1.08E-21	AAEL003566-PA OS=Aedes aegypti GN=AAEL003566 PE=4 SV=1
CCG025243.1	1512	164	43	4.66073841	0.994183797	-2.228974046	Down	5.12E-24	2.06E-23	AAEL009757-PA OS=Aedes aegypti GN=AAEL009757 PE=4 SV=1
CCG005976.1	843	181	43	9.2260047	1.783162398	-2.371267929	Down	2.58E-28	1.08E-27	AAEL001301-PA OS=Aedes aegypti GN=AAEL001301 PE=4 SV=1

CCG015280.1	2310	189	43	3.51570556	0.650738485	-2.433664466	Down	2.20E-30	9.31E-30	AAEL005681-PA OS=Aedes aegypti GN=GPRHIS PE=4 SV=2
CCG001595.2	1908	200	43	4.50416505	0.787843764	-2.515278231	Down	2.86E-33	1.23E-32	AAEL009410-PA OS=Aedes aegypti GN=AAEL009410 PE=4 SV=1
CCG016827.1	3096	251	43	3.48365742	0.485531622	-2.842965595	Down	3.46E-47	1.60E-46	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008110 PE=4 SV=1
CCG011527.1	753	265	43	15.122151	1.996289377	-2.921270591	Down	3.93E-51	1.84E-50	AAEL001235-PA OS=Aedes aegypti GN=AAEL001235 PE=4 SV=1
CCG004982.1	825	339	43	17.6566546	1.822067759	-3.276563505	Down	1.24E-72	6.20E-72	AAEL005386-PA OS=Aedes aegypti GN=AAEL005386 PE=4 SV=1
CCG025361.1	1677	951	43	24.3674524	0.896366071	-4.764723572	Down	3.11E-267	1.93E-266	AAEL009236-PA OS=Aedes aegypti GN=AAEL009236 PE=3 SV=1
CCG026334.1	426	1901	43	191.749919	3.52865235	-5.763964858	Down	0	0	AAEL011504-PA OS=Aedes aegypti GN=AAEL011504 PE=4 SV=1
CCG027096.1	462	0	44	0.001	3.329359693	11.70102903	Up	4.65E-12	1.06E-11	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG023894.1	768	0	44	0.001	2.00281794	10.96781557	Up	4.65E-12	1.06E-11	-
CCG022914.1	1206	0	44	0.001	1.27542635	10.31676388	Up	4.65E-12	1.06E-11	AAEL010627-PA OS=Aedes aegypti GN=AAEL010627 PE=4 SV=1
CCG014180.1	1566	0	44	0.001	0.98222489	9.939909572	Up	4.65E-12	1.06E-11	AAEL002627-PA OS=Aedes aegypti GN=AAEL002627 PE=4 SV=1
CCG025596.1	2079	0	44	0.001	0.739857709	9.531104026	Up	4.65E-12	1.06E-11	AAEL013981-PA OS=Aedes aegypti GN=AAEL013981 PE=4 SV=1
CCG009728.1	4575	0	44	0.001	0.336210749	8.393222041	Up	4.65E-12	1.06E-11	AAEL000067-PA OS=Aedes aegypti GN=AAEL000067 PE=4 SV=1
CCG005063.1	1096	1	44	0.03920596	1.403434469	5.161744822	Up	0.000587	0.00091746	-
CCG016559.1	2223	2	44	0.03865923	0.691931704	4.161744822	Up	9.36E-05	0.00015314	AAEL002752-PA OS=Aedes aegypti GN=AAEL002752 PE=4 SV=1
CCG008314.1	323	2	44	0.26606647	4.762118198	4.161744822	Up	9.36E-05	0.00015315	AAEL000278-PA (Fragment) OS=Aedes aegypti GN=AAEL000278 PE=3 SV=1
CCG017568.1	219	3	44	0.5886265	7.023580721	3.576782322	Up	5.28E-05	8.83E-05	AAEL006867-PA OS=Aedes aegypti GN=AAEL006867 PE=4 SV=1

CCG004534.1	703	4	44	0.24449351	2.188000253	3.161744822	Up	2.96E-05	5.03E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004227 PE=4 SV=1
CCG003736.1	1628	4	44	0.10557674	0.944818291	3.161744822	Up	2.96E-05	5.03E-05	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG020303.1	1299	4	44	0.13231635	1.184114071	3.161744822	Up	2.96E-05	5.03E-05	AAEL007774-PA OS=Aedes aegypti GN=AAEL007774 PE=4 SV=1
CCG015092.1	616	7	44	0.48829244	2.497019769	2.3543899	Up	7.09E-06	1.25E-05	-
CCG010235.2	879	7	44	0.34219356	1.749902364	2.3543899	Up	7.09E-06	1.25E-05	AAEL006902-PA OS=Aedes aegypti GN=AAEL006902 PE=3 SV=1
CCG008092.1	2273	8	44	0.15123532	0.676711033	2.161744822	Up	1.07E-05	1.87E-05	-
CCG001786.1	369	8	44	0.93159316	4.168466607	2.161744822	Up	1.07E-05	1.87E-05	-
CCG000632.1	630	8	44	0.54564742	2.441530441	2.161744822	Up	1.07E-05	1.87E-05	AAEL010719-PA OS=Aedes aegypti GN=AAEL010719 PE=4 SV=1
CCG000703.1	921	8	44	0.37324417	1.670102256	2.161744822	Up	1.07E-05	1.87E-05	-
CCG019126.1	424	8	44	0.81074971	3.627745703	2.161744822	Up	1.07E-05	1.87E-05	AAEL000639-PA OS=Aedes aegypti GN=AAEL000639 PE=3 SV=1
CCG003311.1	1611	9	44	0.24005438	0.954788441	1.991819821	Up	2.48E-05	4.24E-05	AAEL012240-PA (Fragment) OS=Aedes aegypti GN=AAEL012240 PE=4 SV=1
CCG026530.1	525	10	44	0.81847114	2.929836529	1.839816728	Up	6.09E-05	0.00010126	Putative uncharacterized protein OS=Aedes triseriatus PE=2 SV=1
CCG006890.1	2274	10	44	0.18896101	0.676413447	1.839816728	Up	6.09E-05	0.00010126	AAEL007619-PA OS=Aedes aegypti GN=TOLL5A PE=4 SV=1
CCG020659.1	1089	12	44	0.4734957	1.412455627	1.576782322	Up	0.0003151	0.00050057	AAEL012796-PA (Fragment) OS=Aedes aegypti GN=GPRGHP3 PE=3 SV=2
CCG003791.1	2505	75	44	1.286519	0.614037596	-1.067073868	Down	7.14E-05	0.0001183	AAEL001936-PA OS=Aedes aegypti GN=AAEL001936 PE=4 SV=1
CCG006015.1	642	83	44	5.55527722	2.395894358	-1.213294609	Down	3.57E-06	6.41E-06	AAEL014966-PA OS=Aedes aegypti GN=AAEL014966 PE=4 SV=1
CCG012451.1	1071	89	44	3.57078093	1.436194377	-1.313988609	Down	3.21E-07	6.03E-07	Uncharacterized protein OS=Anopheles darlingi GN=AND_05802 PE=4 SV=1

CCG009257.1	1485	89	44	2.57529049	1.035800793	-1.313988609	Down	3.21E-07	6.03E-07	AAEL014343-PA OS=Aedes aegypti GN=AAEL014343 PE=4 SV=1
CCG001782.1	757	97	44	5.5060294	2.031920975	-1.43816802	Down	1.07E-08	2.14E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016435 PE=4 SV=1
CCG021422.1	1416	107	44	3.24700678	1.086274137	-1.579722164	Down	1.17E-10	2.49E-10	-
CCG009480.1	1581	123	44	3.34299643	0.972905868	-1.780769683	Down	5.02E-14	1.69E-13	AAEL004979-PA OS=Aedes aegypti GN=CLIPD2 PE=3 SV=1
CCG000388.1	1434	132	44	3.95537306	1.072638897	-1.882649297	Down	5.05E-16	1.85E-15	AAEL010386-PA (Fragment) OS=Aedes aegypti GN=AAEL010386 PE=3 SV=1
CCG002750.1	873	148	44	7.28467437	1.761929184	-2.047708543	Down	1.00E-19	3.89E-19	AAEL002088-PA OS=Aedes aegypti GN=AAEL002088 PE=4 SV=1
CCG011013.1	615	162	44	11.3188569	2.501079964	-2.17810518	Down	4.19E-23	1.68E-22	AAEL013220-PA OS=Aedes aegypti GN=AAEL013220 PE=4 SV=1
CCG027561.1	357	196	44	23.5912268	4.308583132	-2.452965022	Down	9.86E-32	4.20E-31	AAEL011392-PA OS=Aedes aegypti GN=AAEL011392 PE=4 SV=1
CCG004110.1	1623	210	44	5.55985476	0.947729007	-2.552500695	Down	1.99E-35	8.65E-35	AAEL015586-PA OS=Aedes aegypti GN=AAEL015586 PE=4 SV=1
CCG002641.1	1322	216	44	7.02077358	1.163512994	-2.59314268	Down	4.94E-37	2.17E-36	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010318 PE=4 SV=1
CCG008468.1	411	306	44	31.9920652	3.742491917	-3.09564302	Down	2.79E-62	1.36E-61	Olfant-binding protein 30a OS=Culex quinquefasciatus GN=CpipJ_CPIJ001872 PE=4 SV=1
CCG004016.1	7014	404	44	2.47501751	0.219299141	-3.49646666	Down	1.98E-91	1.03E-90	AAEL002204-PA OS=Aedes aegypti GN=AAEL002204 PE=4 SV=1
CCG007600.1	2472	490	44	8.51746358	0.6222347	-3.774893117	Down	7.13E-118	3.89E-117	AAEL013750-PA OS=Aedes aegypti GN=AAEL013750 PE=4 SV=1
CCG015085.1	1005	818	44	34.9743711	1.53051162	-4.51421221	Down	1.23E-222	7.46E-222	AAEL007500-PA OS=Aedes aegypti GN=AAEL007500 PE=4 SV=1
CCG021117.1	1449	902	44	26.7485857	1.061534974	-4.655238801	Down	4.61E-250	2.84E-249	AAEL012370-PA OS=Aedes aegypti GN=AAEL012370 PE=3 SV=1
CCG015657.1	1131	437487	44	16621.3088	1.360003694	-13.5771358	Down	0	0	AAEL003888-PA OS=Aedes aegypti GN=AAEL003888 PE=4 SV=1

CCG001643.1	940	0	45	0.001	1.673534526	10.7086826	Up	2.56E-12	6.04E-12	AAEL000739-PA OS=Aedes aegypti GN=AAEL000739 PE=4 SV=1
CCG025283.1	3675	5	45	0.05846222	0.428060532	2.872238205	Up	1.61E-05	2.78E-05	Polyprotein OS=Antheraea mylitta PE=4 SV=1
CCG024000.1	734	6	45	0.35125124	2.143218603	2.609203799	Up	8.88E-06	1.56E-05	AAEL000795-PA OS=Aedes aegypti GN=AAEL008795 PE=4 SV=1
CCG015350.1	1888	6	45	0.13655636	0.833221639	2.609203799	Up	8.88E-06	1.56E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC001846 PE=4 SV=1
CCG010915.1	1312	6	45	0.19650793	1.199026261	2.609203799	Up	8.88E-06	1.56E-05	Putative gag-pol protein OS=Drosophila ananassae PE=4 SV=1
CCG010568.1	1476	9	45	0.26201058	1.065801121	2.024241299	Up	1.66E-05	2.86E-05	AAEL014019-PA OS=Aedes aegypti GN=CYP4J17 PE=3 SV=2
CCG015082.1	952	9	45	0.40622648	1.652439553	2.024241299	Up	1.66E-05	2.86E-05	GD14791 OS=Drosophila simulans GN=GD14791 PE=4 SV=1
CCG004054.1	1514	11	45	0.31219754	1.039050499	1.734734682	Up	9.70E-05	0.00015831	Uncharacterized protein OS=Anopheles darlingi GN=AND_05226 PE=4 SV=1
CCG003999.1	2229	11	45	0.21205342	0.705752559	1.734734682	Up	9.70E-05	0.00015832	AAEL010614-PA OS=Aedes aegypti GN=AAEL010614 PE=4 SV=1
CCG009598.1	282	12	45	1.82849935	5.578448421	1.609203799	Up	0.0002172	0.00034839	AAEL010577-PA OS=Aedes aegypti GN=AAEL010577 PE=4 SV=1
CCG011727.1	2078	13	45	0.26881932	0.757036792	1.493726582	Up	0.000458	0.00072143	Uncharacterized protein OS=Anopheles darlingi GN=AND_21982 PE=4 SV=1
CCG016347.1	2913	89	45	1.31284119	0.540035172	-1.281567131	Down	5.39E-07	1.01E-06	AAEL008471-PA OS=Aedes aegypti GN=AAEL008471 PE=4 SV=1
CCG008644.1	1098	89	45	3.48297485	1.432716261	-1.281567131	Down	5.39E-07	1.01E-06	AAEL008024-PA OS=Aedes aegypti GN=AAEL008024 PE=4 SV=1
CCG008245.1	1098	98	45	3.83518578	1.432716261	-1.420543544	Down	1.23E-08	2.46E-08	AAEL009777-PA OS=Aedes aegypti GN=AAEL009777 PE=4 SV=1
CCG018379.1	3558	112	45	1.35261672	0.442136721	-1.613188622	Down	2.16E-11	4.72E-11	AAEL001513-PA OS=Aedes aegypti GN=AAEL001513 PE=4 SV=1

CCG001852.1	663	130	45	8.42543816	2.372733718	-1.828201513	Down	3.09E-15	1.12E-14	AAEL007955-PA (Fragment) OS=Aedes aegypti GN=GSTE8 PE=3 SV=1
CCG013892.1	1002	142	45	6.08952327	1.56998249	-1.955580819	Down	5.96E-18	2.28E-17	AAEL002116-PA OS=Aedes aegypti GN=AAEL002116 PE=4 SV=1
CCG000020.1	168	150	45	38.3658345	9.363824135	-2.03465239	Down	8.09E-20	3.15E-19	AAEL002549-PA OS=Aedes aegypti GN=AAEL002549 PE=4 SV=1
CCG003801.1	1201	169	45	6.04653218	1.309843842	-2.206713136	Down	2.06E-24	8.34E-24	AAEL002149-PA OS=Aedes aegypti GN=AAEL002149 PE=3 SV=1
CCG013634.1	1278	172	45	5.78309417	1.230925238	-2.232098455	Down	3.72E-25	1.51E-24	AAEL004923-PA OS=Aedes aegypti GN=AAEL004923 PE=4 SV=1
CCG011903.1	273	174	45	27.3873034	5.762353314	-2.248777196	Down	1.18E-25	4.83E-25	AAEL001392-PA OS=Aedes aegypti GN=AAEL001392 PE=4 SV=1
CCG002850.1	1335	178	45	5.72929795	1.17836888	-2.281567131	Down	1.17E-26	4.84E-26	AAEL003476-PA OS=Aedes aegypti GN=AAEL003476 PE=4 SV=1
CCG011680.1	1156	199	45	7.39703909	1.360832573	-2.44245832	Down	4.87E-32	2.08E-31	Uncharacterized protein OS=Anopheles darlingi GN=AND_16259 PE=3 SV=1
CCG023957.1	1038	216	45	8.94167888	1.51553223	-2.560721202	Down	1.59E-36	6.97E-36	AAEL009640-PA OS=Aedes aegypti GN=AAEL009640 PE=4 SV=1
CCG016363.1	1107	301	45	11.6837309	1.421068161	-3.039453377	Down	3.35E-60	1.62E-59	AAEL004690-PA OS=Aedes aegypti GN=AAEL004690 PE=4 SV=1
CCG021895.1	13461	521	45	1.66311802	0.1168652	-3.830973262	Down	1.05E-126	5.82E-126	AAEL003155-PA OS=Aedes aegypti GN=AAEL003155 PE=4 SV=1
CCG004173.2	1764	0	46	0.001	0.911610392	9.832273561	Up	1.41E-12	3.42E-12	AAEL007932-PA OS=Aedes aegypti GN=AAEL007932 PE=4 SV=1
CCG004626.1	1344	1	46	0.03197153	1.19648864	5.22587516	Up	0.000587	0.0009174	AAEL008449-PA OS=Aedes aegypti GN=AAEL008449 PE=4 SV=1
CCG017399.1	415	2	46	0.20708306	3.874893329	4.22587516	Up	9.36E-05	0.00015316	-
CCG022013.1	648	2	46	0.13262264	2.481606067	4.22587516	Up	9.36E-05	0.00015317	AAEL011896-PA OS=Aedes aegypti GN=AAEL011896 PE=4 SV=1
CCG004612.1	405	3	46	0.31829433	3.970569707	3.640912659	Up	5.28E-05	8.83E-05	AAEL006515-PA OS=Aedes aegypti GN=AAEL006515 PE=4 SV=1

CCG006731.1	480	3	46	0.26856084	3.350168191	3.640912659	Up	5.28E-05	8.83E-05	AAEL012593-PA (Fragment) OS=Aedes aegypti GN=AAEL012593 PE=4 SV=1
CCG021786.1	945	3	46	0.13641186	1.701672732	3.640912659	Up	5.28E-05	8.83E-05	-
CCG011297.1	1521	3	46	0.08475293	1.057252289	3.640912659	Up	5.28E-05	8.83E-05	AAEL010929-PA OS=Aedes aegypti GN=AAEL010929 PE=3 SV=1
CCG025937.1	693	3	46	0.18601617	2.320462816	3.640912659	Up	5.28E-05	8.83E-05	AAEL011307-PA OS=Aedes aegypti GN=AAEL011307 PE=4 SV=1
CCG004499.1	542	4	46	0.31711981	2.966938619	3.22587516	Up	2.95E-05	5.02E-05	-
CCG016552.1	936	5	46	0.22953918	1.71803497	2.903947065	Up	1.60E-05	2.77E-05	AAEL013440-PA OS=Aedes aegypti GN=AAEL013440 PE=4 SV=1
CCG005924.2	210	5	46	1.02308892	7.657527293	2.903947065	Up	1.60E-05	2.77E-05	AAEL004060-PA OS=Aedes aegypti GN=AAEL004060 PE=4 SV=1
CCG020283.1	1251	6	46	0.20608985	1.285436236	2.640912659	Up	8.69E-06	1.52E-05	AGAP001085-PA (Fragment) OS=Anopheles gambiae GN=AGAP001085 PE=4 SV=1
CCG003556.1	1355	6	46	0.19027189	1.186775448	2.640912659	Up	8.69E-06	1.52E-05	-
CCG022215.1	764	7	46	0.39370176	2.104817711	2.418520238	Up	5.42E-06	9.62E-06	-
CCG014635.1	294	7	46	1.02308892	5.469662352	2.418520238	Up	5.42E-06	9.62E-06	-
CCG005635.1	687	8	46	0.50037537	2.340728867	2.22587516	Up	5.70E-06	1.01E-05	AAEL007040-PA OS=Aedes aegypti GN=AAEL007040 PE=4 SV=1
CCG024026.1	1468	9	46	0.26343843	1.095422842	2.055950158	Up	1.12E-05	1.96E-05	Pol-like protein OS=Aedes aegypti PE=4 SV=1
CCG015973.1	1077	10	46	0.39897618	1.493111171	1.903947065	Up	2.73E-05	4.66E-05	AAEL007298-PA OS=Aedes aegypti GN=AAEL007298 PE=4 SV=1
CCG026661.1	857	10	46	0.50139714	1.876406921	1.903947065	Up	2.73E-05	4.66E-05	AAEL015565-PA OS=Aedes aegypti GN=AAEL015565 PE=4 SV=1
CCG028150.1	1395	10	46	0.30802677	1.152746044	1.903947065	Up	2.73E-05	4.66E-05	AAEL004640-PA OS=Aedes aegypti GN=AAEL004640 PE=4 SV=1
CCG003396.1	705	11	46	0.67044976	2.280965577	1.766443541	Up	6.56E-05	0.00010891	AAEL003906-PA OS=Aedes aegypti GN=AAEL003906 PE=4 SV=1
CCG011886.1	1678	11	46	0.28168479	0.958331783	1.766443541	Up	6.56E-05	0.0001089	AAEL008622-PA OS=Aedes aegypti GN=AAEL008622 PE=4 SV=1
CCG024727.1	540	92	46	7.3207696	2.977927281	-1.297686796	Down	2.66E-07	5.02E-07	AAEL010947-PA OS=Aedes aegypti GN=AAEL010947 PE=4 SV=1

CCG010367.1	1062	94	46	3.80334751	1.514200312	-1.328713692	Down	1.17E-07	2.24E-07	AAEL000558-PA OS=Aedes aegypti GN=AAEL000558 PE=4 SV=1
CCG013635.1	1236	104	46	3.61557638	1.301036191	-1.474564558	Down	1.59E-09	3.26E-09	AAEL004935-PA OS=Aedes aegypti GN=AAEL004935 PE=4 SV=1
CCG004844.1	900	119	46	5.6815538	1.786756368	-1.668942603	Down	1.51E-12	3.64E-12	Glutamate receptor OS=Culex quinquefasciatus GN=CpipJ_CPIJ006667 PE=4 SV=1
CCG017514.1	371	125	46	14.4776734	4.334449411	-1.739909125	Down	8.05E-14	2.58E-13	-
CCG028039.1	1600	132	46	3.54500311	1.005050457	-1.818518959	Down	2.40E-15	8.70E-15	AAEL012390-PA OS=Aedes aegypti GN=AAEL012390 PE=4 SV=1
CCG021355.1	1455	140	46	4.13454491	1.105210125	-1.903407857	Down	3.88E-17	1.47E-16	AAEL007748-PA OS=Aedes aegypti GN=SCRB10 PE=4 SV=2
CCG011562.1	2046	140	46	2.94025555	0.785963212	-1.903407857	Down	3.88E-17	1.47E-16	AAEL004761-PA OS=Aedes aegypti GN=AAEL004761 PE=4 SV=1
CCG017243.1	1824	145	46	3.41590544	0.881623208	-1.95403393	Down	2.79E-18	1.07E-17	AAEL001203-PA OS=Aedes aegypti GN=AAEL001203 PE=4 SV=1
CCG002599.1	825	212	46	11.0419197	1.949188765	-2.502045295	Down	5.77E-35	2.50E-34	Beta-galactosidase OS=Culex quinquefasciatus GN=CpipJ_CPIJ003336 PE=3 SV=1
CCG001145.1	1995	405	46	8.72317921	0.806055505	-3.435902938	Down	2.95E-90	1.54E-89	AAEL000555-PA OS=Aedes aegypti GN=AAEL000555 PE=4 SV=1
CCG002524.1	582	603	46	44.520189	2.763025312	-4.010139032	Down	1.27E-151	7.21E-151	AAEL000419-PA OS=Aedes aegypti GN=AAEL000419 PE=4 SV=1
CCG021133.1	435	0	47	0.001	3.777101169	11.88306371	Up	7.79E-13	1.95E-12	AAEL000566-PA OS=Aedes aegypti GN=AAEL000566 PE=4 SV=1
CCG019402.1	1164	0	47	0.001	1.41154554	10.46305996	Up	7.79E-13	1.95E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_00916 PE=4 SV=1
CCG027664.1	885	3	47	0.14566012	1.856541252	3.671939555	Up	5.28E-05	8.83E-05	AAEL010185-PA OS=Aedes aegypti GN=AAEL010185 PE=4 SV=1
CCG006013.1	564	4	47	0.30474989	2.913189731	3.256902055	Up	2.95E-05	5.02E-05	AAEL013900-PA (Fragment) OS=Aedes aegypti GN=AAEL013900 PE=4 SV=1

CCG010206.1	1173	8	47	0.29305872	1.400715267	2.256902055	Up	4.44E-06	7.90E-06	AAEL001724-PA (Fragment) OS=Aedes aegypti GN=GPRNNB3 PE=4 SV=2
CCG000128.1	1086	8	47	0.3165358	1.512927264	2.256902055	Up	4.44E-06	7.91E-06	Zinc finger protein 307 OS=Culex quinquefasciatus GN=CpipJ_CPIJ015272 PE=4 SV=1
CCG002068.1	632	9	47	0.61191078	2.599745266	2.086977054	Up	7.69E-06	1.36E-05	-
CCG014793.1	666	9	47	0.58067209	2.467025538	2.086977054	Up	7.69E-06	1.36E-05	AAEL008347-PA OS=Aedes aegypti GN=AAEL008347 PE=4 SV=1
CCG026139.1	936	9	47	0.41317053	1.755383556	2.086977054	Up	7.69E-06	1.36E-05	AAEL001481-PA OS=Aedes aegypti GN=AAEL001481 PE=4 SV=1
CCG004468.1	3925	10	47	0.10947703	0.418608665	1.934973961	Up	1.82E-05	3.14E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC001426 PE=4 SV=1
CCG025549.1	780	10	47	0.55089403	2.106460267	1.934973961	Up	1.82E-05	3.14E-05	Transcriptional activator HAPZ OS=Culex quinquefasciatus GN=CpipJ_CPIJ017497 PE=4 SV=1
CCG004064.1	668	13	47	0.83623735	2.459639234	1.556462337	Up	0.0002219	0.00035572	AAEL002668-PA OS=Aedes aegypti GN=AAEL002668 PE=4 SV=1
CCG009375.1	758	14	47	0.79363626	2.167597636	1.449547133	Up	0.0004562	0.00071869	Uncharacterized protein OS=Anopheles darlingi GN=AND_10609 PE=4 SV=1
CCG015778.1	1083	79	47	3.13444971	1.517118198	-1.046878693	Down	5.99E-05	9.96E-05	-
CCG027522.1	528	79	47	6.42918378	3.111816304	-1.046878693	Down	5.99E-05	9.96E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_01498 PE=4 SV=1
CCG005653.1	423	90	47	9.14249673	3.884252975	-1.234951041	Down	9.82E-07	1.81E-06	AAEL012691-PA OS=Aedes aegypti GN=AAEL012691 PE=3 SV=1
CCG025059.1	623	105	47	7.2420901	2.637301779	-1.457343462	Down	1.84E-09	3.77E-09	-
CCG010869.1	3777	109	47	1.24005853	0.435011652	-1.511282269	Down	3.09E-10	6.49E-10	AAEL012104-PA (Fragment) OS=Aedes aegypti GN=AAEL012104 PE=4 SV=1
CCG002404.1	1484	124	47	3.590463	1.107169143	-1.697294255	Down	2.68E-13	7.50E-13	-

CCG023887.1	3484	124	47	1.52934762	0.471595582	-1.697294255	Down	2.68E-13	7.50E-13	GK18646 OS=Drosophila willistoni GN=GK18646 PE=4 SV=1
CCG011273.1	1158	130	47	4.82389076	1.418859247	-1.765465758	Down	1.39E-14	4.96E-14	-
CCG022484.1	540	145	47	11.5381695	3.04266483	-1.923007035	Down	6.36E-18	2.43E-17	AAEL008241-PA OS=Aedes aegypti GN=AAEL008241 PE=4 SV=1
CCG008752.1	4507	154	47	1.46823588	0.364552698	-2.009884485	Down	5.26E-20	2.05E-19	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG001978.1	504	161	47	13.726443	3.259998032	-2.074014823	Down	1.16E-21	4.61E-21	AAEL007682-PA OS=Aedes aegypti GN=AAEL007682 PE=4 SV=1
CCG011216.1	1368	168	47	5.27698495	1.201051907	-2.135415367	Down	2.41E-23	9.66E-23	AAEL009927-PA OS=Aedes aegypti GN=AAEL009927 PE=4 SV=1
CCG015519.1	1128	168	47	6.39974771	1.456594866	-2.135415367	Down	2.41E-23	9.65E-23	AAEL013185-PA OS=Aedes aegypti GN=AAEL013185 PE=4 SV=1
CCG000672.1	1942	306	47	6.77072028	0.846055102	-3.000485787	Down	2.14E-60	1.03E-59	-
CCG027024.1	2418	422	47	7.49926717	0.679503312	-3.464197133	Down	1.17E-94	6.14E-94	AAEL013322-PA OS=Aedes aegypti GN=AAEL013322 PE=4 SV=1
CCG002387.1	3426	575	47	7.21179142	0.479579395	-3.91051609	Down	6.37E-142	3.59E-141	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011995 PE=4 SV=1
CCG015925.1	1581	617	47	16.7693398	1.039240359	-4.012224624	Down	3.65E-155	2.08E-154	AAEL011459-PA OS=Aedes aegypti GN=AAEL011459 PE=4 SV=1
CCG027112.1	1761	652	47	15.9092941	0.933014769	-4.091826099	Down	2.88E-166	1.67E-165	AAEL008943-PA OS=Aedes aegypti GN=AAEL008943 PE=4 SV=1
CCG014071.1	789	701	47	38.177166	2.082432203	-4.196368579	Down	6.69E-182	3.93E-181	Alcohol dehydrogenase z OS=Culex quinquefasciatus GN=CpipJ_CPIJ018908 PE=3 SV=1
CCG011625.1	291	832	47	122.855049	5.646182159	-4.443537663	Down	4.00E-224	2.42E-223	-
CCG020013.1	4959	1085	47	9.40152491	0.331324664	-4.826577272	Down	6.00E-307	3.80E-306	Cadherin OS=Culex quinquefasciatus GN=CpipJ_CPIJ008474 PE=4 SV=1
CCG011849.1	935	1249	47	57.4002123	1.757260971	-5.029655706	Down	0	0	AAEL002512-PA OS=Aedes aegypti GN=AAEL002512 PE=4 SV=1
CCG023728.1	1251	5720	47	196.472328	1.313380502	-7.224897376	Down	0	0	AAEL013367-PA OS=Aedes aegypti GN=AAEL013367 PE=4 SV=1

CCG010580.1	456	0	48	0.001	3.679818608	11.84541894	Up	4.29E-13	1.13E-12	AAEL013835-PA OS=Aedes aegypti GN=AAEL013835 PE=4 SV=1
CCG022113.1	1646	0	48	0.001	1.01943942	9.99356033	Up	4.29E-13	1.13E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG018724.1	1524	1	48	0.02819536	1.101048087	5.287275705	Up	0.000587	0.00091649	AAEL004975-PA OS=Aedes aegypti GN=AAEL004975 PE=4 SV=1
CCG005470.1	319	2	48	0.26940272	5.260179577	4.287275705	Up	9.36E-05	0.00015319	-
CCG026325.1	1680	2	48	0.05115445	0.998807908	4.287275705	Up	9.36E-05	0.0001532	AAEL014180-PA OS=Aedes aegypti GN=AAEL014180 PE=4 SV=1
CCG005657.1	403	2	48	0.2132493	4.163764975	4.287275705	Up	9.36E-05	0.00015318	-
CCG016187.1	783	3	48	0.164635	2.143036124	3.702313204	Up	5.28E-05	8.83E-05	AAEL011537-PA OS=Aedes aegypti GN=AAEL011537 PE=4 SV=1
CCG022585.1	967	3	48	0.13330838	1.735260895	3.702313204	Up	5.28E-05	8.83E-05	-
CCG012222.1	2101	4	48	0.08180816	0.798666009	3.287275705	Up	2.95E-05	5.02E-05	AAEL002840-PA OS=Aedes aegypti GN=AAEL002840 PE=4 SV=1
CCG004439.1	586	5	48	0.36663596	2.863476596	2.96534761	Up	1.60E-05	2.76E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_00155 PE=4 SV=1
CCG005759.1	2193	6	48	0.11756425	0.765160641	2.702313204	Up	8.49E-06	1.49E-05	AAEL007619-PA OS=Aedes aegypti GN=TOLL5A PE=4 SV=1
CCG020383.1	1380	7	48	0.21796242	1.215940062	2.479920782	Up	4.75E-06	8.45E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013304 PE=4 SV=1
CCG027073.1	1055	10	48	0.40729606	1.590518754	1.96534761	Up	1.22E-05	2.13E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002513 PE=4 SV=1
CCG024589.1	1433	12	48	0.3598303	1.170968098	1.702313204	Up	6.95E-05	0.00011535	AAEL014705-PA OS=Aedes aegypti GN=AAEL014705 PE=4 SV=1
CCG016852.1	1686	12	48	0.30583441	0.995253431	1.702313204	Up	6.95E-05	0.00011534	Alkaline phosphatase OS=Aedes aegypti GN=AAEL003309 PE=3 SV=1
CCG024319.1	2364	12	48	0.21812048	0.709812726	1.702313204	Up	6.95E-05	0.00011533	STIP OS=Aedes aegypti PE=2 SV=1

CCG016267.1	1407	14	48	0.42755955	1.192606457	1.479920782	Up	0.0003204	0.00050881	AAEL007131-PA OS=Aedes aegypti GN=AAEL007131 PE=4 SV=1
CCG015212.1	1851	81	48	1.88036116	0.90653554	-1.052574298	Down	4.46E-05	7.51E-05	AAEL003111-PA OS=Aedes aegypti GN=AAEL003111 PE=4 SV=1
CCG026921.1	1431	86	48	2.58238796	1.172604672	-1.13898905	Down	7.28E-06	1.29E-05	AAEL011644-PA OS=Aedes aegypti GN=GPR5HT5B PE=3 SV=1
CCG013182.1	1520	87	48	2.45945192	1.103945582	-1.155667791	Down	5.00E-06	8.90E-06	AAEL003632-PA OS=Aedes aegypti GN=CLIPB39 PE=3 SV=1
CCG002694.1	711	89	48	5.37877128	2.36005244	-1.188457726	Down	2.34E-06	4.22E-06	Protein Wnt OS=Aedes aegypti GN=AAEL008277 PE=3 SV=1
CCG010223.1	864	106	48	5.27174985	1.942126487	-1.44064475	Down	2.13E-09	4.36E-09	AAEL012439-PA OS=Aedes aegypti GN=AAEL012439 PE=4 SV=1
CCG005368.1	2058	110	48	2.29673023	0.815353394	-1.494084009	Down	3.63E-10	7.61E-10	AAEL009939-PA OS=Aedes aegypti GN=AAEL009939 PE=4 SV=1
CCG024731.1	945	112	48	5.09270929	1.775658503	-1.520079218	Down	1.47E-10	3.13E-10	AAEL003294-PA OS=Aedes aegypti GN=AAEL003294 PE=4 SV=1
CCG012702.1	891	117	48	5.6424904	1.88327417	-1.583089015	Down	1.48E-11	3.27E-11	AAEL007953-PA OS=Aedes aegypti GN=AAEL007953 PE=4 SV=1
CCG010498.2	1262	154	48	5.24353338	1.329633348	-1.979510836	Down	1.24E-19	4.80E-19	-
CCG003075.1	2157	189	48	3.76508106	0.777931055	-2.27496672	Down	4.11E-28	1.71E-27	AAEL009045-PA OS=Aedes aegypti GN=AAEL009045 PE=4 SV=1
CCG006479.1	285	221	48	33.3203907	5.887709772	-2.500626855	Down	2.33E-36	1.02E-35	AGAP004476-PA OS=Anopheles gambiae GN=AgaP_AGAP004476 PE=4 SV=4
CCG020420.1	1395	290	48	8.93277637	1.202865437	-2.892633385	Down	3.03E-55	1.44E-54	AAEL004082-PA OS=Aedes aegypti GN=AAEL004082 PE=3 SV=1
CCG006871.1	1260	643	48	21.9282058	1.331743877	-4.041399223	Down	1.66E-162	9.58E-162	AAEL014802-PA OS=Aedes aegypti GN=AAEL014802 PE=4 SV=1
CCG009922.1	3975	892	48	9.64251655	0.422137682	-4.513624195	Down	1.34E-242	8.23E-242	AAEL003262-PA OS=Aedes aegypti GN=AAEL003262 PE=4 SV=1
CCG007672.1	549	2412	48	188.785064	3.056461357	-5.948738487	Down	0	0	Probable GGY family antimicrobial peptide OS=Aedes aegypti PE=4 SV=1
CCG012229.1	489	3	49	0.263618	3.502976609	3.732060547	Up	5.28E-05	8.83E-05	AAEL009296-PA OS=Aedes aegypti GN=AAEL009296 PE=4 SV=1

CCG012174.1	798	4	49	0.21538714	2.146560854	3.317023048	Up	2.95E-05	5.02E-05	AAEL004627-PA OS=Aedes aegypti GN=AAEL004627 PE=4 SV=1
CCG012372.1	1656	5	49	0.12973954	1.034393455	2.995094953	Up	1.60E-05	2.76E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006639 PE=4 SV=1
CCG005139.1	687	6	49	0.37528152	2.493385097	2.732060547	Up	8.45E-06	1.48E-05	AAEL014136-PA OS=Aedes aegypti GN=AAEL014136 PE=4 SV=1
CCG004937.1	1608	8	49	0.21377977	1.065270872	2.317023048	Up	3.10E-06	5.57E-06	Gag-like protein OS=Aedes aegypti PE=4 SV=1
CCG028608.1	1135	8	49	0.30287038	1.509211949	2.317023048	Up	3.10E-06	5.57E-06	GK18652 OS=Drosophila willistoni GN=GK18652 PE=4 SV=1
CCG022792.1	1961	10	49	0.21912154	0.87351125	1.995094953	Up	8.23E-06	1.45E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010720 PE=4 SV=1
CCG010842.1	3012	14	49	0.19972652	0.568710346	1.509668126	Up	0.0002241	0.00035923	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004811 PE=4 SV=1
CCG007551.1	801	14	49	0.75103157	2.138521301	1.509668126	Up	0.0002241	0.00035925	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG026321.1	783	15	49	0.82317499	2.18768271	1.410132452	Up	0.0004506	0.00071015	AAEL002760-PA OS=Aedes aegypti GN=AAEL002760 PE=4 SV=1
CCG008496.1	1977	80	49	1.73878542	0.866441862	-1.004905047	Down	9.44E-05	0.00015421	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC002325 PE=4 SV=1
CCG005090.1	552	85	49	6.61671638	3.103180366	-1.092367888	Down	1.63E-05	2.81E-05	AAEL006454-PA OS=Aedes aegypti GN=AAEL006454 PE=4 SV=1
CCG022227.1	510	89	49	7.49863996	3.358736396	-1.158710383	Down	3.71E-06	6.66E-06	AAEL012245-PA OS=Aedes aegypti GN=AAEL012245 PE=4 SV=1
CCG004469.1	2415	95	49	1.69032082	0.709298369	-1.25283256	Down	3.63E-07	6.81E-07	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC001808 PE=4 SV=1

CCG025598.1	1120	99	49	3.79821761	1.529424609	-1.312333572	Down	7.22E-08	1.39E-07	-
CCG016453.3	1547	112	49	3.11093101	1.107275735	-1.490331874	Down	2.71E-10	5.71E-10	AGAP005610-PA OS=Anopheles gambiae GN=AGAP005610 PE=4 SV=4
CCG019029.1	561	125	49	9.57436154	3.053396723	-1.648761237	Down	6.60E-13	1.68E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009310 PE=4 SV=1
CCG027369.1	1788	132	49	3.17226229	0.958028838	-1.727371071	Down	2.22E-14	7.75E-14	AAEL006156-PA OS=Aedes aegypti GN=AAEL006156 PE=4 SV=1
CCG006042.2	1650	192	49	5.00011457	1.038154886	-2.267939453	Down	1.98E-28	8.25E-28	AAEL007088-PA OS=Aedes aegypti GN=AAEL007088 PE=4 SV=1
CCG001355.1	1683	326	49	8.32331164	1.017798908	-3.031705106	Down	6.69E-65	3.27E-64	AAEL010653-PA OS=Aedes aegypti GN=AAEL010653 PE=4 SV=1
CCG028321.1	1293	674	49	22.3987634	1.324791618	-4.079581733	Down	1.92E-171	1.12E-170	AAEL000044-PA OS=Aedes aegypti GN=CYP325Q1 PE=3 SV=1
CCG027318.1	397	684	49	74.0334974	4.314749526	-4.100829467	Down	1.27E-174	7.40E-174	AAEL002781-PA OS=Aedes aegypti GN=AAEL002781 PE=3 SV=1
CCG006125.1	681	3145	49	198.443194	2.515353248	-6.301821254	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006517 PE=4 SV=1
CCG018685.1	844	4576	49	232.973348	2.029568201	-6.842848289	Down	0	0	AAEL002889-PA OS=Aedes aegypti GN=AAEL002889 PE=4 SV=1
CCG028308.1	1041	14690	49	606.364459	1.645490453	-8.525523728	Down	0	0	AAEL014231-PA (Fragment) OS=Aedes aegypti GN=AAEL014231 PE=4 SV=1
CCG021472.1	804	0	50	0.001	2.174022187	11.08615095	Up	1.30E-13	3.98E-13	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015175 PE=4 SV=1
CCG015992.1	819	0	50	0.001	2.134204931	11.059483	Up	1.30E-13	3.98E-13	AAEL011889-PA OS=Aedes aegypti GN=AAEL011889 PE=3 SV=1
CCG011288.1	1452	0	50	0.001	1.20379741	10.2333769	Up	1.30E-13	3.98E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004879 PE=4 SV=1

CCG020531.1	582	1	50	0.07383116	3.003288382	5.346169394	Up	0.000587	0.00091643	AGAP004106-PA OS=Anopheles gambiae GN=AgaP_AGAP004106 PE=4 SV=5
CCG018339.1	945	1	50	0.04547062	1.849644274	5.346169394	Up	0.000587	0.00091547	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004194 PE=4 SV=1
CCG027277.1	576	1	50	0.07460023	3.034572636	5.346169394	Up	0.000587	0.00091837	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 SV=1
CCG001719.1	687	2	50	0.12509384	2.544270507	4.346169394	Up	9.36E-05	0.00015328	AAEL008482-PA OS=Aedes aegypti GN=AAEL008482 PE=4 SV=1
CCG011508.1	878	3	50	0.14682142	1.990790249	3.761206893	Up	5.28E-05	8.83E-05	Uncharacterized protein (Fragment) OS=Xenopus tropicalis GN=LOC100489570 PE=4 SV=1
CCG020555.1	414	3	50	0.31137489	4.222014103	3.761206893	Up	5.28E-05	8.83E-05	-
CCG025618.1	1380	4	50	0.12454996	1.266604231	3.346169394	Up	2.95E-05	5.02E-05	AAEL004194-PA OS=Aedes aegypti GN=AAEL004194 PE=4 SV=1
CCG009930.1	498	4	50	0.34513843	3.509867146	3.346169394	Up	2.95E-05	5.02E-05	Activated protein kinase c receptor OS=Aedes triseriatus PE=2 SV=1
CCG018405.1	937	6	50	0.27515305	1.865436327	2.761206893	Up	8.42E-06	1.48E-05	AAEL014558-PA OS=Aedes aegypti GN=AAEL014558 PE=4 SV=1
CCG019785.1	1629	6	50	0.1582679	1.072998059	2.761206893	Up	8.42E-06	1.48E-05	AAEL012971-PA OS=Aedes aegypti GN=AAEL012971 PE=4 SV=1
CCG027935.1	821	7	50	0.36636802	2.129005894	2.538814472	Up	4.48E-06	7.98E-06	Retrovirus-related Pol polyprotein from transposon TNT 1-94 OS=Trichinella spiralis GN=Tsp_01017 PE=4 SV=1
CCG005522.1	663	8	50	0.5184885	2.636370797	2.346169394	Up	2.77E-06	4.98E-06	AAEL011476-PA OS=Aedes aegypti GN=AAEL011476 PE=2 SV=1
CCG002000.1	957	8	50	0.35920363	1.826451242	2.346169394	Up	2.77E-06	4.98E-06	-
CCG022513.2	5457	10	50	0.07874241	0.320306732	2.024241299	Up	5.58E-06	9.90E-06	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010686 PE=4 SV=1
CCG000826.1	1540	11	50	0.30692668	1.135008986	1.886737775	Up	1.34E-05	2.33E-05	-

CCG025871.1	1194	11	50	0.39586858	1.463914438	1.886737775	Up	1.34E-05	2.33E-05	AAEL006846-PA OS=Aedes aegypti GN=AAEL006846 PE=4 SV=1
CCG022740.1	474	12	50	1.08784138	3.687581938	1.761206893	Up	3.20E-05	5.42E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002557 PE=4 SV=1
CCG023414.1	359	14	50	1.67569996	4.868840776	1.538814472	Up	0.0001562	0.00025221	-
CCG003347.1	1821	14	50	0.33035491	0.959864821	1.538814472	Up	0.0001562	0.00025219	AAEL005578-PA OS=Aedes aegypti GN=AAEL005578 PE=4 SV=1
CCG013381.1	1431	15	50	0.45041651	1.2214632	1.439278798	Up	0.0003185	0.00050586	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000648 PE=4 SV=1
CCG006911.1	1155	15	50	0.5580485	1.513345315	1.439278798	Up	0.0003185	0.00050589	AAEL005748-PA OS=Aedes aegypti GN=AAEL005748 PE=4 SV=1
CCG011999.1	1953	88	50	1.93616828	0.894989165	-1.113262225	Down	8.41E-06	1.48E-05	AAEL014371-PA OS=Aedes aegypti GN=AAEL014371 PE=3 SV=1
CCG017895.1	578	93	50	6.91381543	3.024072385	-1.192989418	Down	1.29E-06	2.36E-06	-
CCG006615.1	1728	95	50	2.36234073	1.011524212	-1.223686215	Down	5.93E-07	1.10E-06	AAEL011885-PA OS=Aedes aegypti GN=AAEL011885 PE=4 SV=1
CCG022811.1	364	103	50	12.1590183	4.801961095	-1.340331134	Down	2.34E-08	4.61E-08	-
CCG012573.1	1437	125	50	3.73779877	1.216363144	-1.619614891	Down	1.29E-12	3.15E-12	AAEL005409-PA OS=Aedes aegypti GN=AAEL005409 PE=4 SV=1
CCG016153.1	3696	127	50	1.47650333	0.472920411	-1.642515293	Down	5.01E-13	1.30E-12	AAEL011586-PA OS=Aedes aegypti GN=AAEL011586 PE=4 SV=1
CCG025709.1	1305	128	50	4.21465596	1.339397577	-1.653830606	Down	3.11E-13	8.55E-13	AAEL006926-PA OS=Aedes aegypti GN=AAEL006926 PE=4 SV=1
CCG000004.1	1173	133	50	4.8721012	1.490122625	-1.709113042	Down	2.77E-14	9.64E-14	AAEL001262-PA OS=Aedes aegypti GN=AAEL001262 PE=4 SV=1
CCG018224.1	3786	202	50	2.29262715	0.461678246	-2.312042089	Down	1.64E-30	6.94E-30	AAEL001675-PA OS=Aedes aegypti GN=CLIPA10 PE=3 SV=2
CCG004981.1	2151	482	50	9.62873644	0.812605225	-3.566719943	Down	1.21E-110	6.49E-110	AAEL005386-PA OS=Aedes aegypti GN=AAEL005386 PE=4 SV=1
CCG016008.1	711	504	50	30.4595587	2.458387959	-3.63111053	Down	2.20E-117	1.20E-116	AAEL010178-PA OS=Aedes aegypti GN=AAEL010178 PE=4 SV=1

CCG003707.1	4911	524	50	4.58483831	0.35591811	-3.687253608	Down	1.52E-123	8.36E-123	AAEL000307-PA (Fragment) OS=Aedes aegypti GN=AAEL000307 PE=4 SV=1
CCG012503.1	372	644	50	74.3884653	4.698693115	-3.984747485	Down	4.81E-161	2.77E-160	AAEL002725-PA OS=Aedes aegypti GN=AAEL002725 PE=4 SV=1
CCG003288.1	501	0	51	0.001	3.558626977	11.797105	Up	7.17E-14	2.33E-13	AAEL009160-PA OS=Aedes aegypti GN=AAEL009160 PE=4 SV=1
CCG023142.2	1705	0	51	0.001	1.045672795	10.03021577	Up	7.17E-14	2.33E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG000458.2	1500	4	51	0.11458596	1.18858141	3.374738546	Up	2.95E-05	5.02E-05	AAEL005074-PA OS=Aedes aegypti GN=AAEL005074 PE=4 SV=1
CCG026714.1	1515	4	51	0.11345144	1.176813277	3.374738546	Up	2.95E-05	5.02E-05	AAEL003984-PA OS=Aedes aegypti GN=AAEL003984 PE=4 SV=1
CCG022532.1	809	6	51	0.31868777	2.203797423	2.789776045	Up	8.40E-06	1.48E-05	-
CCG017825.1	1377	9	51	0.28084794	1.294750992	2.204813544	Up	2.25E-06	4.06E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016207 PE=4 SV=1
CCG000732.1	909	9	51	0.42544292	1.961355462	2.204813544	Up	2.25E-06	4.06E-06	AAEL013123-PA OS=Aedes aegypti GN=AAEL013123 PE=4 SV=1
CCG021988.1	1261	14	51	0.47706287	1.413855762	1.567383624	Up	0.0001084	0.00017659	-
CCG021174.1	1146	86	51	3.22460487	1.55573483	-1.051526209	Down	2.63E-05	4.51E-05	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012811 PE=4 SV=1
CCG008974.1	658	92	51	6.00792642	2.70953209	-1.14882341	Down	2.98E-06	5.36E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005440 PE=4 SV=1
CCG013968.1	1085	93	51	3.68312011	1.643200106	-1.164420265	Down	2.05E-06	3.72E-06	-
CCG005646.1	1737	94	51	2.32536273	1.026408817	-1.179850306	Down	1.40E-06	2.56E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015620 PE=4 SV=1
CCG018118.1	456	95	51	8.95202804	3.909807271	-1.195117063	Down	9.56E-07	1.76E-06	AAEL003270-PA OS=Aedes aegypti GN=AAEL003270 PE=4 SV=1
CCG011979.1	2980	103	51	1.48519553	0.598279233	-1.311761981	Down	3.97E-08	7.75E-08	-

CCG025821.1	1020	111	51	4.67611818	1.747913839	-1.419677321	Down	1.36E-09	2.80E-09	AAEL010082-PA OS=Aedes aegypti GN=AAEL010082 PE=4 SV=1
CCG022135.1	2808	112	51	1.71389255	0.634925967	-1.432616376	Down	8.83E-10	1.82E-09	AAEL012804-PA (Fragment) OS=Aedes aegypti GN=AAEL012804 PE=4 SV=1
CCG012997.1	573	119	51	8.92390649	3.11146966	-1.520079218	Down	3.94E-11	8.54E-11	AAEL013434-PA OS=Aedes aegypti GN=AAEL013434 PE=4 SV=1
CCG019271.1	801	146	51	7.83218633	2.225807884	-1.815086013	Down	8.89E-17	3.35E-16	AAEL009006-PA OS=Aedes aegypti GN=AAEL009006 PE=4 SV=1
CCG019603.1	1011	148	51	6.29032712	1.763473902	-1.83471482	Down	3.21E-17	1.22E-16	AAEL007364-PA OS=Aedes aegypti GN=AAEL007364 PE=4 SV=1
CCG025996.1	1758	181	51	4.42407393	1.014147961	-2.125107341	Down	6.91E-25	2.81E-24	AAEL003978-PA OS=Aedes aegypti GN=AAEL003978 PE=4 SV=1
CCG017540.1	462	312	51	29.0185221	3.859030553	-2.910663673	Down	1.09E-59	5.24E-59	AAEL001985-PA OS=Aedes aegypti GN=AAEL001985 PE=4 SV=1
CCG000954.2	1710	430	51	10.8052549	1.042615272	-3.373454304	Down	3.66E-94	1.92E-93	AAEL013966-PA OS=Aedes aegypti GN=AAEL013966 PE=4 SV=1
CCG015721.1	567	801	51	60.7032759	3.144395265	-4.270919887	Down	3.57E-210	2.14E-209	AAEL000897-PA OS=Aedes aegypti GN=AAEL000897 PE=4 SV=1
CCG002549.1	561	0	52	0.001	3.24033938	11.66192921	Up	3.95E-14	1.36E-13	AAEL006786-PA OS=Aedes aegypti GN=AAEL006786 PE=3 SV=1
CCG016419.1	689	0	52	0.001	2.638360511	11.365426	Up	3.95E-14	1.36E-13	-
CCG019502.1	876	0	52	0.001	2.075148849	11.01899911	Up	3.95E-14	1.36E-13	AAEL001693-PA OS=Aedes aegypti GN=AAEL001693 PE=3 SV=1
CCG002901.1	1123	0	52	0.001	1.618726974	10.66064396	Up	3.95E-14	1.36E-13	-
CCG009412.1	495	4	52	0.34723018	3.672384631	3.402752922	Up	2.95E-05	5.02E-05	-
CCG006659.1	399	7	52	0.75385499	4.555965895	2.595398	Up	4.38E-06	7.80E-06	AAEL013061-PA OS=Aedes aegypti GN=AAEL013061 PE=4 SV=1
CCG014401.1	658	10	52	0.65303548	2.76266017	2.080824827	Up	2.69E-06	4.85E-06	-
CCG014244.1	1192	11	52	0.39653279	1.525025497	1.943321303	Up	6.04E-06	1.07E-05	Pol polyprotein (Fragment) OS=Anopheles gambiae PE=4 SV=1
CCG003425.1	276	11	52	1.71256189	6.586342001	1.943321303	Up	6.04E-06	1.07E-05	-
CCG025041.1	1098	12	52	0.46961459	1.655583235	1.817790421	Up	1.46E-05	2.53E-05	AAEL008911-PA OS=Aedes aegypti GN=AAEL008911 PE=4 SV=1

CCG007493.1	930	14	52	0.64685622	1.954656336	1.595398	Up	7.50E-05	0.00012414	AAEL003277-PA OS=Aedes aegypti GN=AAEL003277 PE=3 SV=1
CCG007481.1	1131	15	52	0.56989038	1.607277093	1.495862326	Up	0.0001573	0.00025392	AAEL013971-PA OS=Aedes aegypti GN=AAEL013971 PE=4 SV=1
CCG021846.1	996	17	52	0.73341917	1.825130916	1.315290081	Up	0.0006	0.00093369	AAEL012398-PA OS=Aedes aegypti GN=AAEL012398 PE=4 SV=1
CCG003321.1	5139	94	52	0.78598075	0.35373232	-1.15183593	Down	2.22E-06	4.01E-06	AAEL012157-PA (Fragment) OS=Aedes aegypti GN=AAEL012157 PE=4 SV=1
CCG019737.1	1806	96	52	2.28410549	1.006550605	-1.182209579	Down	1.04E-06	1.92E-06	AAEL002343-PA OS=Aedes aegypti GN=AAEL002343 PE=4 SV=1
CCG020563.1	3078	99	52	1.38206749	0.590588172	-1.226603698	Down	3.26E-07	6.13E-07	AAEL012730-PA OS=Aedes aegypti GN=AAEL012730 PE=4 SV=1
CCG001004.1	1138	101	52	3.81365834	1.597390503	-1.255458561	Down	1.48E-07	2.82E-07	AAEL014811-PA OS=Aedes aegypti GN=AAEL014811 PE=4 SV=1
CCG018519.2	723	106	52	6.29985044	2.514288233	-1.325167533	Down	1.96E-08	3.86E-08	AAEL007750-PA OS=Aedes aegypti GN=AAEL007750 PE=4 SV=1
CCG019163.1	459	115	52	10.7658376	3.960414798	-1.442737129	Down	4.26E-10	8.88E-10	Strakimin OS=Culex quinquefasciatus GN=CpipJ_CPIJ004208 PE=4 SV=1
CCG017623.1	1602	126	52	3.37964205	1.134725588	-1.574527002	Down	3.00E-12	7.02E-12	AAEL001312-PA OS=Aedes aegypti GN=CYP9M6 PE=3 SV=1
CCG000554.1	1665	146	52	3.76791667	1.091790025	-1.787071637	Down	1.89E-16	7.10E-16	AAEL002741-PA OS=Aedes aegypti GN=SCRB6 PE=4 SV=2
CCG022866.1	2274	188	52	3.55246707	0.79939771	-2.15183593	Down	3.50E-26	1.44E-25	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003307 PE=4 SV=1
CCG015914.1	1599	195	52	5.24021154	1.136854529	-2.204577392	Down	6.74E-28	2.80E-27	Voltage-dependent para-like sodium channel (Fragment) OS=Aedes albopictus PE=2 SV=1
CCG009459.1	468	199	52	18.2713188	3.884252975	-2.233871699	Down	6.88E-29	2.88E-28	AAEL004753-PA OS=Aedes aegypti GN=AAEL004753 PE=4 SV=1
CCG013641.1	810	211	52	11.1933506	2.244235052	-2.318346267	Down	6.67E-32	2.84E-31	AAEL009974-PA OS=Aedes aegypti GN=AAEL009974 PE=3 SV=1

CCG012835.1	798	308	52	16.5848098	2.277982948	-2.864033619	Down	5.64E-58	2.70E-57	AAEL003532-PA (Fragment) OS=Aedes aegypti GN=AAEL003532 PE=4 SV=1
CCG008912.1	696	515	52	31.7951341	2.611825276	-3.6056757	Down	3.36E-119	1.83E-118	AAEL011752-PA OS=Aedes aegypti GN=GST11 PE=3 SV=2
CCG016354.1	3804	700	52	7.90715411	0.477873394	-4.04845819	Down	5.49E-177	3.21E-176	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018577 PE=4 SV=1
CCG015991.1	819	0	53	0.001	2.262257227	11.14354726	Up	2.18E-14	7.61E-14	AAEL011891-PA OS=Aedes aegypti GN=AAEL011891 PE=3 SV=1
CCG014198.1	1059	0	53	0.001	1.749564371	10.77278003	Up	2.18E-14	7.62E-14	Uncharacterized protein OS=Anopheles darlingi GN=AND_02187 PE=4 SV=1
CCG005014.1	790	1	53	0.05439207	2.345302113	5.430233658	Up	0.000587	0.00091785	-
CCG012020.1	381	2	53	0.22556291	4.862962386	4.430233658	Up	9.36E-05	0.00015379	Uncharacterized protein OS=Anopheles darlingi GN=AND_08690 PE=4 SV=1
CCG003189.1	570	2	53	0.150771	3.250506437	4.430233658	Up	9.36E-05	0.00015394	-
CCG022084.1	1464	2	53	0.05870182	1.265566031	4.430233658	Up	9.36E-05	0.0001538	AAEL007619-PA OS=Aedes aegypti GN=TOLL5A PE=4 SV=1
CCG000729.1	691	6	53	0.37310913	2.681315006	2.845271158	Up	8.39E-06	1.47E-05	-
CCG012064.1	831	6	53	0.31025079	2.229589253	2.845271158	Up	8.39E-06	1.47E-05	AAEL012561-PA OS=Aedes aegypti GN=AAEL012561 PE=4 SV=1
CCG018331.1	1263	7	53	0.23815372	1.466974401	2.622878736	Up	4.35E-06	7.76E-06	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ019142 PE=4 SV=1
CCG018445.1	1890	8	53	0.18188247	0.980311465	2.430233658	Up	2.33E-06	4.21E-06	Acyl-coa dehydrogenase (Fragment) OS=Culex quinquefasciatus GN=CpipJ_CPIJ016452 PE=3 SV=1
CCG005873.1	578	9	53	0.66907891	3.205516728	2.260308657	Up	1.57E-06	2.86E-06	Retrovirus-related Pol polyprotein from transposon TNT 1-94 OS=Trichinella spiralis GN=Tsp_00504 PE=4 SV=1

CCG028153.1	483	10	53	0.88964254	3.836001385	2.108305563	Up	1.94E-06	3.53E-06	AAEL009155-PA OS=Aedes aegypti GN=AAEL009155 PE=4 SV=1
CCG014753.1	1296	10	53	0.33155659	1.429620887	2.108305563	Up	1.94E-06	3.53E-06	AAEL004614-PA OS=Aedes aegypti GN=AAEL004614 PE=4 SV=1
CCG025717.1	843	11	53	0.56069642	2.197851327	1.97080204	Up	4.07E-06	7.29E-06	Moulier or mag4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001640 PE=4 SV=1
CCG019731.1	690	12	53	0.74729973	2.685200969	1.845271158	Up	9.80E-06	1.71E-05	AAEL010667-PA OS=Aedes aegypti GN=AAEL010667 PE=3 SV=1
CCG004378.1	1494	16	53	0.46018457	1.240153058	1.430233658	Up	0.0002226	0.00035683	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG018204.1	1681	17	53	0.43455413	1.10219433	1.342770817	Up	0.0004307	0.00067977	AAEL013982-PA OS=Aedes aegypti GN=AAEL013982 PE=4 SV=1
CCG020580.1	213	17	53	3.42950934	8.698538352	1.342770817	Up	0.0004307	0.00067981	-
CCG009445.2	222	95	53	18.3879495	8.345894905	-1.13962195	Down	2.39E-06	4.32E-06	AAEL010673-PA OS=Aedes aegypti GN=AAEL010673 PE=4 SV=1
CCG006649.1	360	118	53	14.0845241	5.146635191	-1.452409391	Down	2.05E-10	4.33E-10	AAEL005747-PA OS=Aedes aegypti GN=AAEL005747 PE=4 SV=1
CCG003970.1	1518	139	53	3.93464632	1.220545895	-1.688707414	Down	1.23E-14	4.40E-14	AAEL004491-PA OS=Aedes aegypti GN=AAEL004491 PE=4 SV=1
CCG007171.1	2282	148	53	2.7868189	0.811914404	-1.779219707	Down	1.46E-16	5.49E-16	-
CCG017257.1	1563	234	53	6.43308887	1.185405418	-2.440131061	Down	2.35E-37	1.04E-36	AAEL011144-PA OS=Aedes aegypti GN=AAEL011144 PE=3 SV=1
CCG025362.1	1587	264	53	7.1480844	1.167478682	-2.614160461	Down	2.63E-45	1.20E-44	AAEL009237-PA OS=Aedes aegypti GN=AAEL009237 PE=3 SV=1
CCG027139.1	1575	318	53	8.67579404	1.176373758	-2.882649297	Down	3.32E-60	1.60E-59	AAEL005200-PA (Fragment) OS=Aedes aegypti GN=AAEL005200 PE=4 SV=2
CCG009543.1	487	0	54	0.001	3.876277096	11.92045599	Up	1.20E-14	4.28E-14	GL12416 OS=Drosophila persimilis GN=GL12416 PE=3 SV=1
CCG018104.1	468	1	54	0.09181567	4.03364732	5.457200706	Up	0.000587	0.00091757	AAEL009493-PA OS=Aedes aegypti GN=AAEL009493 PE=4 SV=1
CCG009644.1	1275	2	54	0.06740351	1.48058584	4.457200706	Up	9.36E-05	0.00015358	AAEL014550-PA (Fragment) OS=Aedes aegypti GN=AAEL014550 PE=3 SV=1

CCG024719.1	795	2	54	0.10809996	2.37452446	4.457200706	Up	9.36E-05	0.00015331	AAEL006919-PA OS=Aedes aegypti GN=AAEL006919 PE=3 SV=1
CCG015504.1	679	5	54	0.31641925	2.78018696	3.135272611	Up	1.59E-05	2.76E-05	Stromal antigen OS=Culex quinquefasciatus GN=CpipJ_CPIJ009961 PE=4 SV=1
CCG003007.1	1002	6	54	0.2573038	1.883978988	2.872238205	Up	8.38E-06	1.47E-05	AAEL014348-PB OS=Aedes aegypti GN=CASPS8 PE=4 SV=1
CCG019957.1	1656	9	54	0.23353117	1.139943808	2.287275705	Up	1.40E-06	2.55E-06	AAEL011904-PC OS=Aedes aegypti GN=AAEL011904 PE=4 SV=1
CCG021224.1	549	13	54	1.01749827	3.438519027	1.756760988	Up	1.56E-05	2.71E-05	AAEL010722-PA OS=Aedes aegypti GN=AAEL010722 PE=4 SV=1
CCG027783.1	882	14	54	0.68205928	2.14030266	1.649845784	Up	3.55E-05	6.01E-05	AAEL001027-PA OS=Aedes aegypti GN=AAEL001027 PE=4 SV=1
CCG004490.1	1005	15	54	0.64133932	1.87835517	1.55031011	Up	7.65E-05	0.0001266	Putative reverse transcriptase OS=Anopheles gambiae PE=4 SV=1
CCG013308.1	1245	16	54	0.55222149	1.516262607	1.457200706	Up	0.000157	0.00025356	AAEL013617-PA OS=Aedes aegypti GN=AAEL013617 PE=4 SV=1
CCG004544.1	1401	17	54	0.52140292	1.347428227	1.369737865	Up	0.0003079	0.00048958	Uncharacterized protein OS=Anopheles darlingi GN=AND_22260 PE=4 SV=1
CCG023151.1	1410	99	54	3.01702392	1.338827621	-1.172155914	Down	8.39E-07	1.55E-06	AAEL003378-PA (Fragment) OS=Aedes aegypti GN=GPRNNA1 PE=4 SV=2
CCG011778.1	2067	103	54	2.14121077	0.913278638	-1.229299821	Down	1.80E-07	3.41E-07	AAEL011763-PA OS=Aedes aegypti GN=PPO3 PE=4 SV=1
CCG006876.1	989	103	54	4.47510886	1.90874312	-1.229299821	Down	1.80E-07	3.41E-07	AAEL005162-PA OS=Aedes aegypti GN=AAEL005162 PE=3 SV=1
CCG005813.1	576	107	54	7.98222501	3.277338447	-1.28426628	Down	3.67E-08	7.16E-08	-
CCG024738.1	1053	130	54	5.30490551	1.792732142	-1.565167107	Down	1.70E-12	4.08E-12	LINE-1 retrotransposon-like element, putative (Fragment) OS=Aspergillus flavus (strain ATCC 200026 / FGSC A1120 / NRRL 3357 / JCM 12722 / SRRC 167) GN=AFLA_063980 PE=4 SV=1

CCG017760.1	705	134	54	8.16729708	2.677655242	-1.608888484	Down	2.64E-13	7.39E-13	Nuclear hormone receptor OS=Aedes aegypti PE=2 SV=1
CCG011891.1	723	151	54	8.97431525	2.610991626	-1.781204033	Down	6.84E-17	2.58E-16	AAEL000340-PA OS=Aedes aegypti GN=AAEL000340 PE=3 SV=1
CCG003427.1	1128	183	54	6.97115375	1.673534526	-2.058499132	Down	3.45E-24	1.39E-23	AAEL001912-PA OS=Aedes aegypti GN=AAEL001912 PE=4 SV=1
CCG022416.1	1296	203	54	6.73059886	1.456594866	-2.208135211	Down	4.92E-29	2.07E-28	AAEL003953-PA OS=Aedes aegypti GN=AAEL003953 PE=4 SV=1
CCG018423.1	1077	249	54	9.93450689	1.752782679	-2.502801226	Down	8.38E-41	3.75E-40	AAEL007018-PA OS=Aedes aegypti GN=AAEL007018 PE=3 SV=1
CCG008002.1	480	301	54	26.9456044	3.932806137	-2.776418971	Down	6.97E-55	3.31E-54	Hydroxypyruvate isomerase OS=Aedes aegypti GN=AAEL002165 PE=3 SV=1
CCG026199.1	786	548	54	29.9585427	2.401713671	-3.640831377	Down	9.06E-128	5.01E-127	AAEL014108-PA OS=Aedes aegypti GN=AAEL014108 PE=4 SV=1
CCG008058.1	743	697	54	40.309428	2.540709214	-3.98781414	Down	2.95E-174	1.72E-173	Mitochondrial ATP synthase F chain OS=Culex tarsalis PE=4 SV=1
CCG008065.1	351	0	55	0.001	5.477792657	12.41937894	Up	6.66E-15	2.40E-14	-
CCG002773.1	1101	2	55	0.07805583	1.746326269	4.483672917	Up	9.36E-05	0.00015409	AAEL000979-PA OS=Aedes aegypti GN=AAEL000979 PE=4 SV=1
CCG014813.1	1710	2	55	0.050257	1.124389019	4.483672917	Up	9.36E-05	0.00015374	AAEL000747-PA OS=Aedes aegypti GN=AAEL000747 PE=4 SV=1
CCG022296.1	498	3	55	0.25885382	3.86085386	3.898710417	Up	5.28E-05	8.84E-05	AAEL013683-PA OS=Aedes aegypti GN=AAEL013683 PE=4 SV=1
CCG000924.1	606	4	55	0.28362861	3.172780895	3.483672917	Up	2.95E-05	5.02E-05	AAEL013310-PA OS=Aedes aegypti GN=AAEL013310 PE=4 SV=1
CCG007426.1	768	4	55	0.2238007	2.503522425	3.483672917	Up	2.95E-05	5.02E-05	AAEL004104-PA OS=Aedes aegypti GN=AAEL004104 PE=4 SV=1
CCG019498.1	2640	8	55	0.13021132	0.728297433	2.483672917	Up	2.24E-06	4.06E-06	AAEL009404-PA OS=Aedes aegypti GN=AAEL009404 PE=4 SV=1
CCG023319.1	2211	12	55	0.2332143	0.869608875	1.898710417	Up	4.43E-06	7.89E-06	AGAP003323-PA OS=Anopheles gambiae GN=AgaP_AGAP003323 PE=3 SV=5
CCG023855.1	1386	15	55	0.46504042	1.387233205	1.576782322	Up	5.31E-05	8.86E-05	AAEL008547-PA OS=Aedes aegypti GN=AAEL008547 PE=4 SV=1

CCG017887.1	204	18	55	3.79144717	9.4250256	1.313747916	Up	0.0004175	0.00065925	AAEL003160-PA OS=Aedes aegypti GN=AAEL003160 PE=4 SV=1
CCG023924.1	813	90	55	4.75679719	2.364951073	-1.008180179	Down	3.28E-05	5.55E-05	AAEL012645-PA OS=Aedes aegypti GN=AAEL012645 PE=4 SV=1
CCG002727.1	582	99	55	7.30928475	3.303617221	-1.145683703	Down	1.32E-06	2.42E-06	AAEL000419-PA OS=Aedes aegypti GN=AAEL000419 PE=4 SV=1
CCG012015.1	1695	137	55	3.47307	1.134339364	-1.614359166	Down	1.25E-13	3.84E-13	AAEL013023-PA OS=Aedes aegypti GN=AAEL013023 PE=4 SV=1
CCG024659.1	1080	147	55	5.84865832	1.780282613	-1.715999427	Down	1.03E-15	3.74E-15	-
CCG001341.1	1530	151	55	4.24080387	1.25667008	-1.754731822	Down	1.43E-16	5.38E-16	Gooseberry OS=Culex quinquefasciatus GN=CpipJ_CPIJ006496 PE=3 SV=1
CCG021663.1	387	175	55	19.4307585	4.968230549	-1.967538194	Down	6.13E-22	2.43E-21	-
CCG008680.1	3609	183	55	2.17884772	0.532752902	-2.032026921	Down	8.32E-24	3.35E-23	AAEL001494-PA OS=Aedes aegypti GN=AAEL001494 PE=4 SV=1
CCG007634.1	356	192	55	23.1746883	5.400857367	-2.101289583	Down	6.01E-26	2.46E-25	AAEL014300-PA OS=Aedes aegypti GN=AAEL014300 PE=4 SV=1
CCG020101.1	822	222	55	11.6049648	2.339057448	-2.310742949	Down	2.36E-33	1.01E-32	AGAP000697-PA OS=Anopheles gambiae GN=AgaP_AGAP000697 PE=4 SV=4
CCG016863.1	1782	291	55	7.01694319	1.07895916	-2.701202426	Down	1.40E-51	6.57E-51	AAEL007106-PA OS=Aedes aegypti GN=AAEL007106 PE=3 SV=1
CCG007678.2	1629	598	55	15.7740339	1.180297865	-3.740328757	Down	2.07E-142	1.17E-141	AAEL004114-PA OS=Aedes aegypti GN=AAEL004114 PE=4 SV=1
CCG022167.1	4354	902	55	8.9018605	0.441595136	-4.333310706	Down	8.23E-239	5.04E-238	Uncharacterized protein OS=Anopheles darlingi GN=AND_05226 PE=4 SV=1
CCG000788.1	798	30672	55	1651.5886	2.409405041	-9.420961704	Down	0	0	AAEL003049-PA OS=Aedes aegypti GN=AAEL003049 PE=4 SV=1
CCG003899.1	600	0	56	0.001	3.262772499	11.67188268	Up	3.77E-15	1.36E-14	AAEL009434-PA OS=Aedes aegypti GN=AAEL009434 PE=4 SV=1
CCG025524.1	672	0	56	0.001	2.913189731	11.50838395	Up	3.77E-15	1.36E-14	-
CCG002213.1	1042	0	56	0.001	1.878755757	10.87556181	Up	3.77E-15	1.36E-14	-
CCG021054.1	726	2	56	0.11837392	2.696506197	4.509668126	Up	9.36E-05	0.00015398	AAEL012824-PA OS=Aedes aegypti GN=AAEL012824 PE=3 SV=1

CCG010596.1	924	6	56	0.27902425	2.118683441	2.924705625	Up	8.38E-06	1.47E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011306 PE=4 SV=1
CCG021879.1	1305	8	56	0.263416	1.500125287	2.509668126	Up	2.22E-06	4.02E-06	Pol-like protein OS=Aedes aegypti PE=4 SV=1
CCG000256.1	648	9	56	0.59680187	3.021085647	2.339743124	Up	1.22E-06	2.23E-06	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG012764.1	564	10	56	0.76187473	3.471034573	2.187740031	Up	9.21E-07	1.70E-06	AAEL010373-PA OS=Aedes aegypti GN=AAEL010373 PE=4 SV=1
CCG027969.1	822	11	56	0.57502078	2.381585766	2.050236507	Up	1.34E-06	2.46E-06	AAEL013971-PA OS=Aedes aegypti GN=AAEL013971 PE=4 SV=1
CCG005786.1	4401	12	56	0.11716356	0.444822427	1.924705625	Up	2.98E-06	5.36E-06	AAEL008598-PA OS=Aedes aegypti GN=AAEL008598 PE=4 SV=1
CCG020250.1	857	14	56	0.70195599	2.284321469	1.702313204	Up	1.66E-05	2.86E-05	-
CCG002997.1	2589	92	56	1.52692761	0.756146581	-1.01389383	Down	2.45E-05	4.21E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001896 PE=4 SV=1
CCG022273.1	1711	97	56	2.43603989	1.144163354	-1.090244716	Down	4.25E-06	7.61E-06	AAEL003213-PA OS=Aedes aegypti GN=AAEL003213 PE=4 SV=1
CCG025616.1	834	99	56	5.10072389	2.347318344	-1.119688494	Down	2.06E-06	3.74E-06	FAD oxidoreductase OS=Culex quinquefasciatus GN=CpipJ_CPIJ016820 PE=4 SV=1
CCG006861.1	892	102	56	4.91357952	2.194690022	-1.162757216	Down	6.76E-07	1.25E-06	Reverse transcriptase OS=Anopheles gambiae PE=4 SV=1
CCG017654.1	822	118	56	6.16840473	2.381585766	-1.372974923	Down	1.12E-09	2.31E-09	AAEL000024-PA OS=Aedes aegypti GN=AAEL000024 PE=4 SV=1
CCG020123.1	5202	121	56	0.99948825	0.376329008	-1.409195111	Down	3.13E-10	6.57E-10	AAEL005155-PA OS=Aedes aegypti GN=AAEL005155 PE=4 SV=1
CCG013397.1	1843	146	56	3.40400502	1.062215681	-1.680156433	Down	3.37E-15	1.22E-14	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC006955 PE=4 SV=1

CCG024702.1	1620	257	56	6.81680358	1.208434259	-2.495956423	Down	6.23E-42	2.80E-41	AAEL005533-PA (Fragment) OS=Aedes aegypti GN=AAEL005533 PE=4 SV=1
CCG010215.1	4608	339	56	3.1611849	0.424840169	-2.895473337	Down	2.30E-64	1.12E-63	AAEL011867-PA OS=Aedes aegypti GN=AAEL011867 PE=4 SV=1
CCG013320.1	916	387	56	18.1542438	2.137187226	-3.08652163	Down	4.22E-78	2.13E-77	Copia OS=Drosophila simulans GN=GIP PE=4 SV=1
CCG017548.1	1491	654	56	18.8478916	1.312986921	-3.8434787	Down	5.10E-159	2.93E-158	Amino acid transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ007170 PE=4 SV=1
CCG018253.1	2148	886	56	17.7240153	0.911388966	-4.281494763	Down	1.10E-232	6.69E-232	AAEL009203-PA OS=Aedes aegypti GN=AAEL009203 PE=3 SV=1
CCG010539.1	486	0	57	0.001	4.100044807	12.00142396	Up	2.22E-15	8.04E-15	tRNA-dihydrouridine synthase OS=Aedes aegypti GN=AAEL005898 PE=3 SV=1
CCG023824.1	522	0	57	0.001	3.817283096	11.89833047	Up	2.22E-15	8.05E-15	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 SV=1
CCG011106.1	2979	0	57	0.001	0.668889485	9.385624056	Up	2.22E-15	8.05E-15	AAEL003766-PA OS=Aedes aegypti GN=AAEL003766 PE=4 SV=1
CCG005446.1	750	1	57	0.05729298	2.656829035	5.535203218	Up	0.000587	0.00091694	AAEL002960-PA OS=Aedes aegypti GN=AAEL002960 PE=4 SV=1
CCG016055.1	968	2	57	0.08878044	2.05849357	4.535203218	Up	9.36E-05	0.00015378	AAEL013652-PA OS=Aedes aegypti GN=AAEL013652 PE=3 SV=1
CCG024059.1	2064	2	57	0.04163734	0.965417527	4.535203218	Up	9.36E-05	0.00015323	AAEL012229-PA OS=Aedes aegypti GN=AAEL012229 PE=4 SV=1
CCG003644.1	1182	2	57	0.07270683	1.685805225	4.535203218	Up	9.36E-05	0.00015335	AAEL014020-PA OS=Aedes aegypti GN=AAEL014020 PE=4 SV=1
CCG026649.1	636	3	57	0.20268743	3.133053107	3.950240717	Up	5.28E-05	8.86E-05	-
CCG019966.1	1032	4	57	0.16654936	1.930835054	3.535203218	Up	2.95E-05	5.03E-05	AAEL009977-PA OS=Aedes aegypti GN=AAEL009977 PE=4 SV=1
CCG018647.1	2949	4	57	0.0582838	0.675694058	3.535203218	Up	2.95E-05	5.04E-05	GM18987 OS=Drosophila sechellia GN=GM18987 PE=4 SV=1
CCG006413.1	342	5	57	0.62821249	5.826379462	3.213275123	Up	1.59E-05	2.76E-05	-
CCG026976.1	630	8	57	0.54564742	3.162891708	2.535203218	Up	2.21E-06	3.99E-06	-

CCG001282.1	892	8	57	0.38537879	2.233880915	2.535203218	Up	2.21E-06	3.99E-06	AAEL010264-PA OS=Aedes aegypti GN=AAEL010264 PE=4 SV=1
CCG004161.1	707	8	57	0.48622048	2.818418354	2.535203218	Up	2.21E-06	3.99E-06	-
CCG027064.1	309	9	57	1.25154567	6.448614162	2.365278217	Up	1.17E-06	2.15E-06	AAEL003080-PA OS=Aedes aegypti GN=AAEL003080 PE=4 SV=1
CCG017318.1	579	14	57	1.03899186	3.441488387	1.727848296	Up	1.13E-05	1.97E-05	AAEL001815-PA OS=Aedes aegypti GN=AAEL001815 PE=4 SV=1
CCG012507.1	937	14	57	0.64202378	2.126597413	1.727848296	Up	1.13E-05	1.97E-05	AAEL001621-PA OS=Aedes aegypti GN=AAEL001621 PE=4 SV=1
CCG005667.1	717	18	57	1.07873811	2.779109869	1.365278217	Up	0.0002148	0.00034468	AAEL014714-PA OS=Aedes aegypti GN=AAEL014714 PE=4 SV=1
CCG003820.1	2436	18	57	0.31751035	0.817989235	1.365278217	Up	0.0002148	0.00034473	AAEL001230-PA OS=Aedes aegypti GN=AAEL001230 PE=4 SV=1
CCG017370.1	294	18	57	2.63080008	6.777625089	1.365278217	Up	0.0002148	0.0003447	AAEL010419-PA OS=Aedes aegypti GN=AAEL010419 PE=4 SV=1
CCG020227.1	702	18	57	1.10178807	2.838492558	1.365278217	Up	0.0002148	0.00034466	AAEL000465-PA OS=Aedes aegypti GN=AAEL000465 PE=4 SV=1
CCG016688.1	1335	96	57	3.08995844	1.492600581	-1.049759283	Down	9.20E-06	1.61E-05	Oxysterol-binding protein OS=Aedes aegypti GN=AAEL012056 PE=3 SV=1
CCG003717.2	2284	97	57	1.82489679	0.872426347	-1.064709624	Down	6.47E-06	1.15E-05	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012788 PE=4 SV=1
CCG027446.1	736	98	57	5.72151358	2.707366544	-1.079506626	Down	4.54E-06	8.08E-06	-
CCG012133.1	513	109	57	9.13002158	3.884252975	-1.232981107	Down	7.36E-08	1.42E-07	AAEL001068-PA OS=Aedes aegypti GN=AAEL001068 PE=4 SV=1
CCG013073.1	351	120	57	14.6905076	5.676985117	-1.371687378	Down	8.36E-10	1.73E-09	AAEL006255-PA OS=Aedes aegypti GN=AAEL006255 PE=4 SV=1
CCG007590.1	1461	131	57	3.85286464	1.363875274	-1.498219784	Down	6.98E-12	1.57E-11	AAEL005176-PA OS=Aedes aegypti GN=AAEL005176 PE=4 SV=1
CCG004206.1	1179	154	57	5.61267102	1.690094806	-1.731583323	Down	1.38E-16	5.21E-16	AAEL000196-PA OS=Aedes aegypti GN=AAEL000196 PE=3 SV=1

CCG011023.1	1923	163	57	3.6422604	1.036204772	-1.813524936	Down	1.55E-18	5.95E-18	Chondroitin beta-1,4-N-acetyl galactosaminyl transferase 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ019283 PE=4 SV=1
CCG009743.1	1368	166	57	5.2141637	1.456594866	-1.839836213	Down	3.36E-19	1.30E-18	AAEL001404-PA OS=Aedes aegypti GN=AAEL001404 PE=4 SV=1
CCG010875.1	2316	167	57	3.09842214	0.860372097	-1.848501074	Down	2.02E-19	7.82E-19	-
CCG025508.1	405	205	57	21.7501126	4.920053768	-2.144276882	Down	2.75E-28	1.14E-27	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008280 PE=4 SV=1
CCG022867.1	2331	274	57	5.05092548	0.854835597	-2.562828865	Down	6.09E-46	2.79E-45	AAEL012615-PA OS=Aedes aegypti GN=AAEL012615 PE=4 SV=1
CCG006671.1	1395	316	57	9.73364598	1.428402707	-2.76857753	Down	2.31E-57	1.10E-56	Beta-galactosidase OS=Aedes aegypti GN=AAEL004575 PE=3 SV=1
CCG004920.1	773	754	57	41.9135574	2.577777201	-4.023217495	Down	1.28E-189	7.58E-189	AAEL002192-PA OS=Aedes aegypti GN=AAEL002192 PE=4 SV=1
CCG011624.1	567	1649	57	124.968417	3.51432412	-5.152172465	Down	0	0	-
CCG000283.1	756	5204	57	295.786374	2.63574309	-6.810202029	Down	0	0	AAEL011798-PA OS=Aedes aegypti GN=AAEL011798 PE=4 SV=1
CCG009600.1	429	0	58	0.001	4.726293829	12.20649361	Up	1.33E-15	4.84E-15	AAEL005568-PA OS=Aedes aegypti GN=AAEL005568 PE=4 SV=1
CCG025780.1	1164	1	58	0.03691558	1.741907262	5.560294199	Up	0.000587	0.00091552	AAEL000130-PA OS=Aedes aegypti GN=AAEL000130 PE=4 SV=1
CCG022185.1	5617	2	58	0.01529989	0.360972059	4.560294199	Up	9.36E-05	0.000154	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018804 PE=4 SV=1
CCG002202.2	1522	2	58	0.05646483	1.332181375	4.560294199	Up	9.36E-05	0.00015341	AAEL015565-PA OS=Aedes aegypti GN=AAEL015565 PE=4 SV=1
CCG013902.1	423	3	58	0.30474989	4.793333458	3.975331698	Up	5.28E-05	8.86E-05	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG012569.1	653	3	58	0.19741073	3.105023052	3.975331698	Up	5.28E-05	8.85E-05	AAEL002948-PA OS=Aedes aegypti GN=AAEL002948 PE=4 SV=1
CCG022226.1	834	3	58	0.15456739	2.431151142	3.975331698	Up	5.28E-05	8.84E-05	AAEL009244-PA OS=Aedes aegypti GN=AAEL009244 PE=3 SV=1

CCG012241.1	414	8	58	0.83033304	4.897536359	2.560294199	Up	2.20E-06	3.98E-06	AAEL006570-PA OS=Aedes aegypti GN=AAEL006570 PE=4 SV=1
CCG007635.1	665	12	58	0.77539371	3.048992561	1.975331698	Up	1.38E-06	2.52E-06	-
CCG018032.1	2425	14	58	0.24807269	0.836115486	1.752939277	Up	7.67E-06	1.35E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004811 PE=4 SV=1
CCG001438.1	1409	14	58	0.42695265	1.439020619	1.752939277	Up	7.67E-06	1.35E-05	AAEL003264-PA OS=Aedes aegypti GN=AAEL003264 PE=4 SV=1
CCG004575.1	1155	16	58	0.59525173	1.755480565	1.560294199	Up	3.76E-05	6.34E-05	AAEL007864-PA OS=Aedes aegypti GN=AAEL007864 PE=4 SV=1
CCG000715.1	2064	16	58	0.33309872	0.982354677	1.560294199	Up	3.76E-05	6.34E-05	Starlet OS=Culex quinquefasciatus
CCG025851.1	4728	100	58	0.90883533	0.428845189	-1.083561991	Down	3.38E-06	6.07E-06	AAEL005045-PA OS=Aedes aegypti GN=AAEL005045 PE=3 SV=1
CCG017641.1	1896	118	58	2.67427673	1.069398762	-1.32234885	Down	3.29E-09	6.69E-09	Transporter OS=Aedes aegypti GN=AAEL000298 PE=3 SV=1
CCG022791.1	1131	123	58	4.67310111	1.792732142	-1.382220306	Down	4.08E-10	8.51E-10	AAEL000577-PA OS=Aedes aegypti GN=AAEL000577 PE=4 SV=1
CCG018921.1	1365	133	58	4.18679465	1.485406632	-1.494988237	Down	5.23E-12	1.19E-11	AAEL000912-PA OS=Aedes aegypti GN=AAEL000912 PE=4 SV=1
CCG014904.1	498	182	58	15.7037986	4.071445889	-1.947500441	Down	1.82E-22	7.26E-22	AAEL013220-PA OS=Aedes aegypti GN=AAEL013220 PE=4 SV=1
CCG027033.1	592	188	58	13.6457941	3.424966305	-1.994294653	Down	7.46E-24	3.00E-23	AAEL006341-PA OS=Aedes aegypti GN=AAEL006341 PE=4 SV=1
CCG012716.1	2058	225	58	4.69785728	0.985218685	-2.253486992	Down	8.50E-33	3.63E-32	AAEL013496-PA OS=Aedes aegypti GN=PPO8 PE=4 SV=1
CCG007640.1	471	352	58	32.1132624	4.304840876	-2.89913742	Down	7.43E-67	3.65E-66	AGAP005599-PA (Fragment) OS=Anopheles gambiae GN=AGAP005599 PE=4 SV=3
CCG006358.1	879	391	58	19.1139548	2.30668948	-3.050730598	Down	5.45E-78	2.75E-77	AAEL008701-PA OS=Aedes aegypti GN=AAEL008701 PE=4 SV=1
CCG015353.1	1353	482	58	15.3077695	1.49858097	-3.352595137	Down	9.44E-105	5.03E-104	AAEL011440-PA OS=Aedes aegypti GN=AAEL011440 PE=4 SV=1

CCG011850.1	1137	895	58	33.8240215	1.783271814	-4.245449673	Down	1.17E-233	7.13E-233	Lipase OS=Aedes aegypti GN=AAEL015326 PE=3 SV=1
CCG013958.1	2970	2804	58	40.5680592	0.682686886	-5.892976435	Down	0	0	AAEL001953-PA OS=Aedes aegypti GN=AAEL001953 PE=4 SV=1
CCG016691.1	1269	4590	58	155.422444	1.597777819	-6.603984239	Down	0	0	AGAP009759-PA OS=Anopheles gambiae GN=AgaP_AGAP009759 PE=4 SV=4
CCG021211.1	288	0	59	0.001	7.161591422	12.8060645	Up	8.88E-16	3.24E-15	AAEL004817-PA OS=Aedes aegypti GN=AAEL004817 PE=4 SV=1
CCG004836.1	774	0	59	0.001	2.664778204	11.37979974	Up	8.88E-16	3.24E-15	AAEL010110-PA OS=Aedes aegypti GN=AAEL010110 PE=4 SV=1
CCG019747.1	1059	0	59	0.001	1.947628262	10.92750263	Up	8.88E-16	3.24E-15	Uncharacterized protein OS=Anopheles darlingi GN=AND_02187 PE=4 SV=1
CCG007299.1	1080	2	59	0.07957358	1.909757713	4.584956253	Up	9.36E-05	0.00015388	AAEL004561-PA OS=Aedes aegypti GN=AAEL004561 PE=4 SV=1
CCG003987.1	861	6	59	0.29944066	2.395514901	2.999993752	Up	8.38E-06	1.47E-05	AAEL004449-PA OS=Aedes aegypti GN=AAEL004449 PE=4 SV=1
CCG005930.1	801	6	59	0.32187067	2.574954219	2.999993752	Up	8.38E-06	1.47E-05	AAEL007204-PA OS=Aedes aegypti GN=AAEL007204 PE=4 SV=1
CCG009535.1	681	6	59	0.37858797	3.028690645	2.999993752	Up	8.38E-06	1.47E-05	AAEL006501-PA OS=Aedes aegypti GN=AAEL006501 PE=4 SV=1
CCG004461.1	1158	7	59	0.25974796	1.781121183	2.777601331	Up	4.31E-06	7.69E-06	AAEL014577-PA OS=Aedes aegypti GN=AAEL014577 PE=4 SV=1
CCG004167.1	589	7	59	0.51067596	3.501762869	2.777601331	Up	4.31E-06	7.69E-06	AAEL012468-PA OS=Aedes aegypti GN=AAEL012468 PE=4 SV=1
CCG012102.1	1257	7	59	0.23929049	1.640841949	2.777601331	Up	4.31E-06	7.69E-06	Uncharacterized protein OS=Anopheles darlingi GN=AND_20812 PE=4 SV=1
CCG019814.1	789	10	59	0.54461007	2.614117021	2.263028158	Up	6.43E-07	1.19E-06	-
CCG000485.1	3156	11	59	0.14976777	0.653529255	2.125524635	Up	5.64E-07	1.05E-06	AGAP002803-PA OS=Anopheles gambiae GN=AgaP_AGAP002803 PE=4 SV=2

CCG000705.1	216	11	59	2.18827352	9.548788563	2.125524635	Up	5.64E-07	1.05E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007491 PE=4 SV=1
CCG026464.1	6514	13	59	0.08575477	0.316631613	1.884516535	Up	2.18E-06	3.97E-06	Putative uncharacterized protein OS=Drosophila simulans GN=pol PE=4 SV=1
CCG009603.1	2195	17	59	0.33279521	0.939652998	1.497493412	Up	5.44E-05	9.08E-05	AAEL005592-PA OS=Aedes aegypti GN=AAEL005592 PE=4 SV=1
CCG003012.1	363	17	59	2.01235672	5.681923773	1.497493412	Up	5.44E-05	9.08E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002500 PE=4 SV=1
CCG014693.3	2985	19	59	0.2735092	0.690967615	1.33702874	Up	0.0002092	0.00033598	AAEL009780-PA OS=Aedes aegypti GN=AAEL009780 PE=3 SV=1
CCG002761.1	865	138	59	6.85528714	2.384437375	-1.523568204	Down	1.02E-12	2.51E-12	AAEL014552-PA OS=Aedes aegypti GN=AAEL014552 PE=4 SV=1
CCG015372.2	1390	138	59	4.26605998	1.483840525	-1.523568204	Down	1.02E-12	2.51E-12	AAEL006581-PA OS=Aedes aegypti GN=AAEL006581 PE=4 SV=1
CCG014200.1	708	145	59	8.80029876	2.913189731	-1.594952837	Down	4.08E-14	1.40E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003070 PE=4 SV=1
CCG018890.1	921	147	59	6.85836155	2.239455298	-1.614716092	Down	1.60E-14	5.64E-14	AAEL011416-PA OS=Aedes aegypti GN=AAEL011416 PE=4 SV=1
CCG007207.1	1587	154	59	4.1697159	1.29964608	-1.681830288	Down	5.69E-16	2.08E-15	AAEL009774-PA OS=Aedes aegypti GN=AAEL009774 PE=3 SV=1
CCG018196.1	1611	251	59	6.69485002	1.2802845	-2.386587301	Down	6.06E-39	2.69E-38	AAEL002666-PA OS=Aedes aegypti GN=AAEL002666 PE=4 SV=1
CCG015445.1	2613	257	59	4.22626169	0.789337286	-2.420668296	Down	1.71E-40	7.64E-40	AAEL002754-PA OS=Aedes aegypti GN=AAEL002754 PE=4 SV=1
CCG010415.1	1473	374	59	10.9101702	1.400229687	-2.961938207	Down	1.69E-72	8.43E-72	AAEL004923-PA OS=Aedes aegypti GN=AAEL004923 PE=4 SV=1
CCG022345.1	1131	377	59	14.3232449	1.823641317	-2.97346446	Down	2.36E-73	1.18E-72	AAEL001951-PA OS=Aedes aegypti GN=AAEL001951 PE=2 SV=1
CCG017574.1	603	640	59	45.6063518	3.420461575	-3.736971842	Down	3.53E-152	2.01E-151	AAEL005392-PA OS=Aedes aegypti GN=AAEL005392 PE=4 SV=1

CCG013937.1	1293	3	60	0.09969776	1.622193818	4.024241299	Up	5.28E-05	8.86E-05	AAEL002128-PA OS=Aedes aegypti GN=AAEL002128 PE=3 SV=1
CCG022132.1	1686	7	60	0.17840341	1.244066789	2.801848877	Up	4.31E-06	7.69E-06	Alkaline phosphatase OS=Aedes aegypti GN=AAEL003309 PE=3 SV=1
CCG027600.1	1341	8	60	0.25634443	1.564128715	2.609203799	Up	2.19E-06	3.96E-06	AAEL006871-PA OS=Aedes aegypti GN=AAEL006871 PE=4 SV=1
CCG006002.1	420	10	60	1.02308892	4.994039539	2.287275705	Up	6.07E-07	1.13E-06	Insulin-like peptide 2 OS=Aedes aegypti PE=2 SV=2
CCG007241.1	729	12	60	0.70732073	2.877224426	2.024241299	Up	6.69E-07	1.24E-06	AAEL014795-PA OS=Aedes aegypti GN=AAEL014795 PE=4 SV=1
CCG011341.1	459	14	60	1.31062371	4.569709382	1.801848877	Up	3.51E-06	6.29E-06	AAEL010292-PA (Fragment) OS=Aedes aegypti GN=AAEL010292 PE=4 SV=1
CCG023223.1	589	15	60	1.09430564	3.561114782	1.702313204	Up	8.13E-06	1.44E-05	-
CCG016120.1	1135	16	60	0.60574075	1.848014631	1.609203799	Up	1.80E-05	3.10E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000390 PE=4 SV=1
CCG017734.1	2172	19	60	0.37588626	0.965698253	1.361276286	Up	0.00015	0.00024252	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG028489.1	1376	21	60	0.6557881	1.524343464	1.216886377	Up	0.0005067	0.00079557	-
CCG006411.3	2331	104	60	1.9171396	0.899826944	-1.091235919	Down	1.88E-06	3.43E-06	AAEL002732-PA (Fragment) OS=Aedes aegypti GN=AAEL002732 PE=4 SV=1
CCG020480.1	642	122	60	8.1655882	3.26712867	-1.321533538	Down	1.82E-09	3.71E-09	AAEL011962-PA OS=Aedes aegypti GN=AAEL011962 PE=4 SV=1
CCG019799.1	1377	135	60	4.21271908	1.523236461	-1.467611798	Down	7.12E-12	1.60E-11	AAEL008262-PA (Fragment) OS=Aedes aegypti GN=AAEL008262 PE=4 SV=1
CCG027704.2	762	136	60	7.66913899	2.752620218	-1.478259042	Down	4.57E-12	1.05E-11	AAEL006641-PA OS=Aedes aegypti GN=AAEL006641 PE=4 SV=1
CCG008075.1	489	150	60	13.1809002	4.289359113	-1.619614891	Down	7.56E-15	2.71E-14	AAEL004665-PA OS=Aedes aegypti GN=AAEL004665 PE=4 SV=1

CCG000173.1	591	152	60	11.0514377	3.549063632	-1.638723714	Down	2.94E-15	1.06E-14	AAEL013424-PA OS=Aedes aegypti GN=AAEL013424 PE=4 SV=1
CCG007529.1	2213	153	60	2.97079503	0.947806871	-1.648184043	Down	1.82E-15	6.62E-15	Gap-Pol polyprotein OS=Schistosoma japonicum PE=4 SV=1
CCG017424.1	432	158	60	15.7157826	4.855316218	-1.694576949	Down	1.65E-16	6.20E-16	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016740 PE=4 SV=1
CCG009208.1	1902	160	60	3.61469902	1.102784756	-1.712724295	Down	6.23E-17	2.35E-16	AAEL002576-PA OS=Aedes aegypti GN=AAEL002576 PE=3 SV=1
CCG017725.1	660	178	60	11.5888072	3.178025161	-1.866529632	Down	7.39E-21	2.91E-20	AAEL006267-PA OS=Aedes aegypti GN=AAEL006267 PE=3 SV=1
CCG001026.1	414	191	60	19.8242012	5.066416924	-1.968225029	Down	8.16E-24	3.28E-23	AAEL014303-PA OS=Aedes aegypti GN=AAEL014303 PE=4 SV=1
CCG021317.1	12798	1499	60	5.03294516	0.163892531	-4.940580869	Down	0	0	AAEL008036-PA OS=Aedes aegypti GN=AAEL008036 PE=4 SV=1
CCG008382.1	756	0	61	0.001	2.820707517	11.46184136	Up	4.44E-16	1.66E-15	AAEL001674-PA OS=Aedes aegypti GN=AAEL001674 PE=3 SV=1
CCG021075.1	453	1	61	0.09485593	4.707405923	5.633050541	Up	0.000587	0.00091603	-
CCG007143.1	1635	2	61	0.05256237	1.304253751	4.633050541	Up	9.36E-05	0.00015363	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018804 PE=4 SV=1
CCG002930.1	204	2	61	0.42127191	10.45321021	4.633050541	Up	9.36E-05	0.00015389	AAEL003533-PA OS=Aedes aegypti GN=AAEL003533 PE=4 SV=1
CCG001060.1	585	4	61	0.29381015	3.645222022	3.633050541	Up	2.95E-05	5.03E-05	-
CCG009488.1	593	5	61	0.36230805	3.596045334	3.311122446	Up	1.59E-05	2.76E-05	-
CCG017195.1	546	6	61	0.47219489	3.905595024	3.048088041	Up	8.38E-06	1.47E-05	AAEL002330-PA OS=Aedes aegypti GN=AAEL002330 PE=4 SV=1
CCG005870.1	862	7	61	0.34894216	2.473845572	2.825695619	Up	4.31E-06	7.69E-06	-
CCG012718.1	2061	8	61	0.16679179	1.034670006	2.633050541	Up	2.19E-06	3.96E-06	AAEL013492-PA OS=Aedes aegypti GN=PPO5 PE=4 SV=1
CCG001451.1	564	15	61	1.14281209	3.780948374	1.726159946	Up	5.54E-06	9.83E-06	AAEL008356-PA OS=Aedes aegypti GN=AAEL008356 PE=4 SV=1
CCG001669.1	1362	15	61	0.47323496	1.565679062	1.726159946	Up	5.54E-06	9.83E-06	AAEL005519-PA OS=Aedes aegypti GN=AAEL005519 PE=4 SV=1

CCG013346.1	562	18	61	1.37625485	3.794403707	1.46312554	Up	5.45E-05	9.09E-05	AAEL007574-PA OS=Aedes aegypti GN=AAEL007574 PE=4 SV=1
CCG017664.1	1569	21	61	0.57512073	1.359117198	1.240733119	Up	0.0003704	0.00058625	AAEL004191-PA OS=Aedes aegypti GN=AAEL004191 PE=4 SV=1
CCG008543.1	1606	114	61	3.05015551	1.327805033	-1.199839473	Down	7.18E-08	1.38E-07	AAEL000486-PA OS=Aedes aegypti GN=AAEL006486 PE=4 SV=1
CCG008826.1	2007	176	61	3.76814813	1.062508661	-1.826381077	Down	4.47E-20	1.75E-19	AAEL012949-PA (Fragment) OS=Aedes aegypti GN=AAEL012949 PE=4 SV=1
CCG015084.1	1035	177	61	7.34844737	2.060342882	-1.834555009	Down	2.69E-20	1.05E-19	AAEL007497-PA OS=Aedes aegypti GN=AAEL007497 PE=4 SV=1
CCG023271.1	1526	178	61	5.01219709	1.397414733	-1.84268289	Down	1.61E-20	6.33E-20	AAEL000438-PA OS=Aedes aegypti GN=AAEL000438 PE=4 SV=1
CCG016996.1	633	185	61	12.5582953	3.368807082	-1.898330919	Down	4.36E-22	1.73E-21	Uncharacterized protein OS=Anopheles darlingi GN=AND_05893 PE=4 SV=1
CCG000766.1	240	195	61	34.9129094	8.88522868	-1.974279772	Down	2.25E-24	9.07E-24	AAEL014137-PA OS=Aedes aegypti GN=CLIPB25 PE=3 SV=1
CCG014951.1	1761	219	61	5.34376597	1.210934062	-2.141736518	Down	4.55E-30	1.92E-29	AAEL009856-PA OS=Aedes aegypti GN=AAEL009856 PE=4 SV=1
CCG010864.1	1132	318	61	12.0710032	1.883794066	-2.679832414	Down	9.88E-56	4.70E-55	AAEL012101-PA OS=Aedes aegypti GN=AAEL012101 PE=3 SV=1
CCG012590.1	873	332	61	16.3412966	2.442674551	-2.74198889	Down	1.44E-59	6.94E-59	AAEL003656-PA OS=Aedes aegypti GN=AAEL003656 PE=4 SV=1
CCG011257.1	1287	3634	61	121.330237	1.656919101	-6.194292163	Down	0	0	AAEL003425-PB (Fragment) OS=Aedes aegypti GN=AAEL003425 PE=4 SV=1
CCG027173.2	1340	4	62	0.12826786	1.617472507	3.656509514	Up	2.95E-05	5.03E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC001807 PE=4 SV=1
CCG020150.1	1935	4	62	0.08882632	1.12011016	3.656509514	Up	2.95E-05	5.04E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015738 PE=4 SV=1
CCG004769.1	297	4	62	0.57871696	7.297687407	3.656509514	Up	2.95E-05	5.04E-05	-

CCG023961.1	914	6	62	0.28207703	2.37134919	3.071547013	Up	8.38E-06	1.47E-05	-
CCG027325.1	768	6	62	0.33570105	2.822152552	3.071547013	Up	8.38E-06	1.47E-05	-
CCG000891.1	777	8	62	0.44241683	2.789463526	2.656509514	Up	2.19E-06	3.96E-06	Arsin OS=Culex quinquefasciatus GN=CpipJ_CPIJ008260 PE=4 SV=1
CCG003620.1	450	10	62	0.95488299	4.816473689	2.334581419	Up	5.69E-07	1.06E-06	-
CCG026422.2	1011	13	62	0.55252873	2.143831019	1.956069796	Up	6.83E-07	1.27E-06	AAEL006948-PA (Fragment) OS=Aedes aegypti GN=AAEL006948 PE=4 SV=1
CCG021735.1	1680	13	62	0.3325039	1.290126881	1.956069796	Up	6.83E-07	1.27E-06	-
CCG024836.1	1179	14	62	0.51024282	1.838348736	1.849154592	Up	1.60E-06	2.92E-06	AAEL014577-PA OS=Aedes aegypti GN=AAEL014577 PE=4 SV=1
CCG009165.1	1176	14	62	0.51154446	1.843038401	1.849154592	Up	1.60E-06	2.92E-06	S-methyl-5-uraciladenosine phosphorylase OS=Culex quinquefasciatus GN=CpipJ_CPIJ002278 PE=3 SV=1
CCG009575.1	1210	14	62	0.49717048	1.791250545	1.849154592	Up	1.60E-06	2.92E-06	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC002325 PE=4 SV=1
CCG009908.1	1341	14	62	0.44860275	1.616266338	1.849154592	Up	1.60E-06	2.92E-06	AAEL009050-PA OS=Aedes aegypti GN=AAEL009050 PE=4 SV=1
CCG018330.1	669	15	62	0.96344696	3.239780508	1.749618919	Up	3.76E-06	6.74E-06	AAEL002806-PA OS=Aedes aegypti GN=AAEL002806 PE=4 SV=1
CCG014474.1	1360	15	62	0.4739309	1.593686147	1.749618919	Up	3.76E-06	6.74E-06	-
CCG012895.1	659	16	62	1.04327125	3.288942579	1.656509514	Up	8.53E-06	1.50E-05	Predicted protein OS=Ajellomyces capsulata (strain NAm1 / WU24) GN=HCAG_05350 PE=4 SV=1
CCG025904.1	270	16	62	2.54635464	8.027456148	1.656509514	Up	8.53E-06	1.50E-05	AAEL004552-PA OS=Aedes aegypti GN=AAEL004552 PE=4 SV=1
CCG028149.1	529	17	62	1.38087994	4.097189338	1.569046673	Up	1.85E-05	3.18E-05	AAEL004617-PA OS=Aedes aegypti GN=AAEL004617 PE=4 SV=1
CCG025897.1	1239	18	62	0.62425765	1.749324584	1.486584513	Up	3.83E-05	6.47E-05	Argininosuccinate synthase OS=Culex quinquefasciatus GN=CpipJ_CPIJ012882 PE=3 SV=1

CCG027089.2	5015	19	62	0.1627966	0.432186074	1.408582001	Up	7.62E-05	0.00012619	Polyprotein OS=Antheraea mylitta PE=4 SV=1
CCG011652.1	1196	20	62	0.71855743	1.812218361	1.334581419	Up	0.000146	0.00023647	Pol-like protein OS=Aedes aegypti PE=4 SV=1
CCG025238.1	1200	105	62	3.75985178	1.806177633	-1.057736003	Down	3.04E-06	5.47E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011883 PE=4 SV=1
CCG006035.2	10573	107	62	0.43485875	0.204995097	-1.084957472	Down	1.50E-06	2.74E-06	AGAP013037-PA OS=Anopheles gambiae GN=AgaP_AGAP013037 PE=4 SV=1
CCG014345.2	621	113	62	7.81896942	3.490198325	-1.163669448	Down	1.67E-07	3.16E-07	AAEL006947-PA (Fragment) OS=Aedes aegypti GN=GPRTAK2 PE=3 SV=2
CCG026266.1	858	128	62	6.4104033	2.526122564	-1.343490486	Down	4.40E-10	9.17E-10	AAEL010102-PA OS=Aedes aegypti GN=AAEL010102 PE=4 SV=1
CCG011863.1	1290	129	62	4.29697346	1.68016524	-1.354717741	Down	2.90E-10	6.10E-10	AAEL014370-PA (Fragment) OS=Aedes aegypti GN=AAEL014370 PE=4 SV=1
CCG021738.1	3384	137	62	1.73961396	0.640488522	-1.441522569	Down	9.55E-12	2.13E-11	AAEL009813-PA OS=Aedes aegypti GN=AAEL009813 PE=4 SV=1
CCG008603.1	1566	193	62	5.29575912	1.384044163	-1.935947523	Down	1.49E-23	5.98E-23	AAEL001241-PA OS=Aedes aegypti GN=AAEL001241 PE=4 SV=1
CCG020911.1	1863	238	62	5.48942396	1.163399442	-2.238308249	Down	2.54E-34	1.10E-33	AAEL004002-PA OS=Aedes aegypti GN=AAEL004002 PE=3 SV=1
CCG027689.1	1332	243	62	7.83907321	1.627187057	-2.268302989	Down	1.44E-35	6.25E-35	AAEL001722-PA OS=Aedes aegypti GN=AAEL001722 PE=4 SV=2
CCG024889.1	1515	250	62	7.09071528	1.430635749	-2.30927477	Down	2.49E-37	1.09E-36	AAEL005983-PA OS=Aedes aegypti GN=AAEL005983 PE=4 SV=1
CCG001210.1	875	306	62	15.02713	2.477043611	-2.600878328	Down	5.84E-52	2.75E-51	-
CCG001796.1	363	0	63	0.001	6.067138944	12.56680064	Up	4.44E-16	1.65E-15	-
CCG007624.1	642	2	63	0.1338621	3.430485104	4.679593127	Up	9.36E-05	0.00015383	AAEL010375-PA OS=Aedes aegypti GN=AAEL010375 PE=4 SV=1
CCG005589.1	1556	4	63	0.11046204	1.415405808	3.679593127	Up	2.95E-05	5.03E-05	AAEL014132-PA OS=Aedes aegypti GN=AAEL014132 PE=4 SV=1

CCG005374.1	648	6	63	0.39786791	3.398721353	3.094630627	Up	8.38E-06	1.47E-05	AAEL011990-PA (Fragment) OS=Aedes aegypti GN=AAEL011990 PE=4 SV=1
CCG001601.1	1251	10	63	0.34348309	1.760488758	2.357665032	Up	5.59E-07	1.04E-06	AAEL011805-PA (Fragment) OS=Aedes aegypti GN=AAEL011805 PE=4 SV=1
CCG001956.1	246	12	63	2.09608462	8.952729417	2.094630627	Up	2.81E-07	5.28E-07	AAEL007576-PA OS=Aedes aegypti GN=AAEL007576 PE=4 SV=1
CCG013893.1	1318	13	63	0.42382895	1.67099502	1.979153409	Up	4.73E-07	8.83E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001645 PE=4 SV=1
CCG011982.1	1295	14	63	0.46453767	1.700672924	1.872238205	Up	1.08E-06	1.99E-06	AAEL004639-PA OS=Aedes aegypti GN=AAEL004639 PE=4 SV=1
CCG000401.2	1425	15	63	0.452313	1.545523815	1.772702532	Up	2.55E-06	4.60E-06	AAEL004576-PA OS=Aedes aegypti GN=AAEL004576 PE=4 SV=1
CCG002159.1	623	18	63	1.24150116	3.53510664	1.509668126	Up	2.69E-05	4.60E-05	AAEL011939-PA OS=Aedes aegypti GN=AAEL011939 PE=4 SV=1
CCG003386.1	960	19	63	0.85044266	2.294136913	1.431665614	Up	5.41E-05	9.03E-05	Annexin OS=Aedes aegypti GN=AAEL005417 PE=3 SV=1
CCG016939.1	435	20	63	1.97561998	5.062922843	1.357665032	Up	0.0001048	0.00017082	AAEL002191-PA OS=Aedes aegypti GN=AAEL002191 PE=4 SV=1
CCG010728.1	885	21	63	1.01962082	2.488555296	1.287275705	Up	0.0001957	0.00031482	AAEL012641-PA OS=Aedes aegypti GN=AAEL012641 PE=4 SV=1
CCG021376.1	1668	23	63	0.59250833	1.320366569	1.156031171	Up	0.0006178	0.00096051	AAEL010649-PA OS=Aedes aegypti GN=AAEL010649 PE=4 SV=1
CCG018400.1	931	23	63	1.06155091	2.365597676	1.156031171	Up	0.0006178	0.00096045	-
CCG010230.1	1610	119	63	3.17602386	1.367932569	-1.215224636	Down	2.70E-08	5.30E-08	AAEL000522-PA OS=Aedes aegypti GN=AAEL000522 PE=4 SV=1
CCG000461.1	267	137	63	22.048141	8.24858216	-1.418438956	Down	1.70E-11	3.73E-11	AAEL011100-PA OS=Aedes aegypti GN=AAEL011100 PE=4 SV=1
CCG010145.1	1926	140	63	3.12344904	1.143495035	-1.44968989	Down	4.61E-12	1.05E-11	AAEL011030-PA OS=Aedes aegypti GN=AAEL011030 PE=4 SV=1
CCG009795.1	766	156	63	8.75101645	2.875158533	-1.605809092	Down	3.28E-15	1.19E-14	-
CCG007325.1	504	203	63	17.3072542	4.369784597	-1.98574279	Down	1.70E-25	6.95E-25	AAEL009620-PA OS=Aedes aegypti GN=AAEL009620 PE=4 SV=1

CCG018412.1	1171	213	63	7.81601492	1.880761261	-2.055116493	Down	7.67E-28	3.18E-27	AAEL004509-PA OS=Aedes aegypti GN=AAEL004509 PE=4 SV=1
CCG006155.1	642	217	63	14.524038	3.430485104	-2.081958105	Down	8.58E-29	3.59E-28	AAEL014966-PA OS=Aedes aegypti GN=AAEL014966 PE=4 SV=1
CCG008460.1	1641	223	63	5.83927533	1.342091064	-2.121306773	Down	3.12E-30	1.31E-29	AAEL001936-PA OS=Aedes aegypti GN=AAEL001936 PE=4 SV=1
CCG013755.1	3517	945	63	11.545749	0.626207403	-4.204577392	Down	6.07E-245	3.73E-244	AAEL011196-PA OS=Aedes aegypti GN=AAEL011196 PE=4 SV=1
CCG014538.1	846	1	64	0.05079165	2.64459777	5.702313204	Up	0.000587	0.00091569	AAEL006359-PA OS=Aedes aegypti GN=AAEL006359 PE=4 SV=1
CCG009804.1	873	1	64	0.04922077	2.562806086	5.702313204	Up	0.000587	0.000917	AAEL005972-PA OS=Aedes aegypti GN=AAEL005972 PE=4 SV=1
CCG017412.1	878	3	64	0.14682142	2.548211519	4.117350703	Up	5.28E-05	8.84E-05	-
CCG020857.1	1872	3	64	0.06886175	1.195154761	4.117350703	Up	5.28E-05	8.85E-05	AAEL013428-PA OS=Aedes aegypti GN=AAEL013428 PE=4 SV=1
CCG002474.1	1107	4	64	0.15526553	2.021074719	3.702313204	Up	2.95E-05	5.04E-05	AAEL009769-PA OS=Aedes aegypti GN=AAEL009769 PE=4 SV=1
CCG000897.1	348	5	64	0.61738124	6.429108372	3.380385109	Up	1.59E-05	2.76E-05	Ccicbpin A OS=Culex quinquefasciatus GN=CpipJ_CPIJ010699 PE=4 SV=1
CCG016632.1	1193	5	64	0.18009109	1.875381151	3.380385109	Up	1.59E-05	2.76E-05	AAEL003811-PA OS=Aedes aegypti GN=AAEL003811 PE=4 SV=1
CCG020417.1	681	6	64	0.37858797	3.285359344	3.117350703	Up	8.38E-06	1.47E-05	AAEL004465-PA OS=Aedes aegypti GN=AAEL004465 PE=3 SV=1
CCG004364.1	1107	11	64	0.4269802	2.021074719	2.242881585	Up	3.00E-07	5.64E-07	AAEL008798-PA OS=Aedes aegypti GN=AAEL008798 PE=4 SV=1
CCG008036.1	513	12	64	1.00513999	4.361266498	2.117350703	Up	2.28E-07	4.32E-07	CCA-adding enzyme OS=Culex quinquefasciatus GN=CpipJ_CPIJ013325 PE=3 SV=1
CCG000109.1	496	12	64	1.03959035	4.51074539	2.117350703	Up	2.28E-07	4.32E-07	AGAP003907-PA OS=Anopheles gambiae GN=AGAP003907 PE=4 SV=2
CCG027826.1	681	13	64	0.82027394	3.285359344	2.001873486	Up	3.33E-07	6.25E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011583 PE=4 SV=1

CCG004654.1	345	16	64	1.99279929	6.485013662	1.702313204	Up	3.99E-06	7.15E-06	AAEL012940-PA OS=Aedes aegypti GN=AAEL012940 PE=4 SV=1
CCG023526.2	759	18	64	1.01904509	2.947733483	1.532388202	Up	1.88E-05	3.24E-05	AAEL008673-PA OS=Aedes aegypti GN=AAEL008673 PE=3 SV=1
CCG013604.1	2438	19	64	0.33487488	0.917690613	1.45438569	Up	3.83E-05	6.46E-05	Retrovirus-related Pol polyprotein from transposon TNT 1-94 OS=Trichinella spiralis GN=Tsp_01017 PE=4 SV=1
CCG022285.1	646	19	64	1.26381572	3.46335869	1.45438569	Up	3.83E-05	6.46E-05	-
CCG023989.1	678	23	64	1.45767536	3.299896332	1.178751248	Up	0.0004566	0.00071918	AAEL012676-PA OS=Aedes aegypti GN=AAEL012676 PE=4 SV=1
CCG025122.1	1102	109	64	4.25018246	2.030244749	-1.065871121	Down	1.70E-06	3.10E-06	AAEL007107-PA OS=Aedes aegypti GN=AAEL007107 PE=3 SV=1
CCG027940.1	1719	111	64	2.77466	1.301529793	-1.092102663	Down	8.35E-07	1.54E-06	AAEL002880-PA OS=Aedes aegypti GN=AAEL002880 PE=3 SV=1
CCG025828.1	756	114	64	6.47956316	2.959430838	-1.13057681	Down	2.81E-07	5.29E-07	Uncharacterized protein OS=Anopheles darlingi GN=AND_05262 PE=4 SV=1
CCG017284.1	891	120	64	5.78716964	2.511032226	-1.204577392	Down	2.95E-08	5.79E-08	AAEL012777-PA OS=Aedes aegypti GN=AAEL012777 PE=3 SV=1
CCG002333.1	2103	123	64	2.51320844	1.06387528	-1.240201302	Down	9.20E-09	1.84E-08	AAEL000134-PA OS=Aedes aegypti GN=AAEL000134 PE=4 SV=1
CCG011879.1	1243	124	64	4.28660265	1.799943454	-1.251883107	Down	6.21E-09	1.25E-08	AAEL007050-PA OS=Aedes aegypti GN=AAEL007050 PE=3 SV=1
CCG002960.1	1051	133	64	5.43765433	2.12876281	-1.352969232	Down	1.61E-10	3.41E-10	-
CCG012634.1	8082	146	64	0.77624118	0.27682872	-1.487511355	Down	5.91E-13	1.52E-12	AAEL011166-PA (Fragment) OS=Aedes aegypti GN=AAEL011166 PE=4 SV=1
CCG017691.1	1323	179	64	5.81374338	1.691103336	-1.781502573	Down	9.48E-20	3.69E-19	AAEL012767-PA OS=Aedes aegypti GN=AAEL012767 PE=4 SV=1
CCG027513.1	702	184	64	11.2627225	3.187079364	-1.821248752	Down	7.64E-21	3.00E-20	AAEL005218-PA OS=Aedes aegypti GN=AAEL005218 PE=4 SV=1
CCG010745.1	1143	184	64	6.91726261	1.957418822	-1.821248752	Down	7.64E-21	3.00E-20	AGAP005258-PA OS=Anopheles gambiae GN=AGAP005258 PE=4 SV=4

CCG005998.1	3432	208	64	2.60422634	0.651902597	-1.998126514	Down	2.74E-26	1.13E-25	AAEL001566-PA OS=Aedes aegypti GN=AAEL001566 PE=3 SV=1
CCG028472.1	2021	211	64	4.48620188	1.107040927	-2.018785985	Down	5.45E-27	2.25E-26	AAEL005003-PA OS=Aedes aegypti GN=AAEL005003 PE=3 SV=1
CCG010082.1	735	263	64	15.3755649	3.043986005	-2.336605785	Down	9.81E-40	4.37E-39	AAEL000820-PA OS=Aedes aegypti GN=AAEL008826 PE=4 SV=1
CCG019258.1	1494	739	64	21.254775	1.497543316	-3.82711735	Down	7.49E-179	4.38E-178	AAEL003568-PA OS=Aedes aegypti GN=AAEL003568 PE=4 SV=1
CCG007963.1	2517	896	64	15.2963378	0.888887451	-4.105041718	Down	2.27E-228	1.38E-227	AAEL000927-PA OS=Aedes aegypti GN=AAEL006927 PE=4 SV=1
CCG008028.1	2383	2	65	0.03606356	0.953540911	4.724681017	Up	9.36E-05	0.00015371	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG020266.1	357	6	65	0.72218041	6.364952353	3.139718516	Up	8.38E-06	1.47E-05	AAEL008355-PA OS=Aedes aegypti GN=AAEL008355 PE=4 SV=1
CCG001700.1	738	8	65	0.46579658	3.078981017	2.724681017	Up	2.19E-06	3.96E-06	GK19301 OS=Drosophila willistoni GN=GK19301 PE=4 SV=1
CCG025343.1	436	11	65	1.08409881	5.211669702	2.265249398	Up	2.88E-07	5.42E-07	-
CCG001614.1	942	11	65	0.50176972	2.412195319	2.265249398	Up	2.88E-07	5.42E-07	Elongation factor-1 beta,delta OS=Culex quinquefasciatus GN=CpipJ_CPIJ004079 PE=3 SV=1
CCG023417.1	261	19	65	3.12806497	8.706084254	1.476753503	Up	2.70E-05	4.61E-05	Putative uncharacterized protein GLEAN_07454 OS=Tribolium castaneum GN=GLEAN_07454 PE=4 SV=1
CCG001599.1	885	22	65	1.06817419	2.567557051	1.265249398	Up	0.0001882	0.00030292	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006742 PE=4 SV=1
CCG021559.1	1080	24	65	0.95488299	2.103970361	1.139718516	Up	0.0005823	0.00091159	AAEL007779-PA OS=Aedes aegypti GN=AAEL007779 PE=4 SV=1
CCG019566.1	1920	120	65	2.68560841	1.183483328	-1.182209579	Down	4.70E-08	9.15E-08	AAEL004391-PA OS=Aedes aegypti GN=AAEL004391 PE=4 SV=1
CCG000349.1	1293	129	65	4.28700369	1.757376636	-1.286546239	Down	1.40E-09	2.87E-09	AAEL005959-PA OS=Aedes aegypti GN=AAEL005959 PE=4 SV=1
CCG024112.1	1083	131	65	5.1976318	2.098142189	-1.308741985	Down	6.21E-10	1.29E-09	AAEL014334-PA OS=Aedes aegypti GN=AAEL014334 PE=4 SV=1

CCG028350.1	1233	135	65	4.70471547	1.842893747	-1.35213458	Down	1.20E-10	2.55E-10	AAEL007224-PA OS=Aedes aegypti GN=AAEL007224 PE=4 SV=1
CCG014643.1	1094	143	65	5.61670206	2.077045695	-1.43519032	Down	3.98E-12	9.19E-12	AAEL000123-PA OS=Aedes aegypti GN=AAEL000123 PE=4 SV=1
CCG025510.1	3849	145	65	1.61876111	0.590358013	-1.455228073	Down	1.66E-12	3.99E-12	AAEL012395-PA OS=Aedes aegypti GN=AAEL012395 PE=3 SV=1
CCG018455.1	2064	148	65	3.08116314	1.100914724	-1.484772349	Down	4.43E-13	1.16E-12	AAEL002294-PA (Fragment) OS=Aedes aegypti GN=AAEL002294 PE=4 SV=1
CCG008866.1	816	152	65	8.00416625	2.784666655	-1.523246497	Down	7.38E-14	2.39E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015436 PE=4 SV=1
CCG023602.2	450	169	65	16.1375226	5.049528867	-1.676198419	Down	2.63E-17	9.99E-17	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016740 PE=4 SV=1
CCG016390.1	2031	221	65	4.67568259	1.118802555	-2.063221543	Down	5.72E-29	2.40E-28	AAEL013801-PA OS=Aedes aegypti GN=AAEL013801 PE=4 SV=1
CCG011870.1	2006	262	65	5.61219864	1.132745758	-2.308741985	Down	4.92E-39	2.19E-38	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001097 PE=4 SV=1
CCG016253.1	1929	274	65	6.10352892	1.177961633	-2.373351066	Down	4.32E-42	1.95E-41	AAEL013371-PA OS=Aedes aegypti GN=AAEL013371 PE=4 SV=1
CCG017894.1	1091	425	65	16.7388975	2.082757095	-3.006638014	Down	2.10E-83	1.07E-82	Pyrroline-5-carboxylate reductase OS=Aedes aegypti PE=2 SV=1
CCG009782.1	537	17096	65	1367.98991	4.231448771	-8.336690176	Down	0	0	AAEL006575-PA OS=Aedes aegypti GN=AAEL006575 PE=4 SV=1
CCG024723.1	768	6	66	0.33570105	3.00422691	3.161744822	Up	8.38E-06	1.47E-05	Putative reverse transcriptase OS=Anopheles gambiae PE=4 SV=1
CCG020806.1	330	8	66	1.04169054	6.991655354	2.746707323	Up	2.19E-06	3.96E-06	AAEL001177-PA OS=Aedes aegypti GN=AAEL001177 PE=4 SV=1
CCG027916.1	1511	8	66	0.22750356	1.526966424	2.746707323	Up	2.19E-06	3.96E-06	Pao retrotransposon peptidase superfamily (Fragment) OS=Trichinella spiralis GN=Tsp_12554 PE=4 SV=1

CCG014402.1	790	9	66	0.48952862	2.920564895	2.576782322	Up	1.10E-06	2.01E-06	-
CCG012089.1	1572	10	66	0.27334437	1.46771391	2.424779228	Up	5.47E-07	1.02E-06	AAEL003746-PA OS=Aedes aegypti GN=CYP9AE1 PE=3 SV=1
CCG018680.1	552	10	66	0.77843722	4.179793962	2.424779228	Up	5.47E-07	1.02E-06	AAEL014169-PA OS=Aedes aegypti GN=AAEL014169 PE=4 SV=1
CCG001235.1	777	11	66	0.60832314	2.969428915	2.287275705	Up	2.81E-07	5.28E-07	AAEL009682-PA OS=Aedes aegypti GN=AAEL009682 PE=3 SV=1
CCG005223.1	1114	12	66	0.46286967	2.071136685	2.161744822	Up	1.72E-07	3.26E-07	-
CCG012517.1	453	13	66	1.23312704	5.093258867	2.046267605	Up	1.80E-07	3.41E-07	AAEL008821-PA OS=Aedes aegypti GN=AAEL008821 PE=4 SV=1
CCG020826.1	1506	17	66	0.48505013	1.532036034	1.659244482	Up	4.19E-06	7.50E-06	AAEL010908-PA OS=Aedes aegypti GN=AAEL010908 PE=4 SV=1
CCG020794.1	1251	21	66	0.72131449	1.844321556	1.3543899	Up	7.33E-05	0.00012138	AAEL005490-PA OS=Aedes aegypti GN=AAEL005490 PE=4 SV=1
CCG006925.1	1608	24	66	0.64133932	1.434854644	1.161744822	Up	0.0004318	0.00068138	AAEL006823-PA OS=Aedes aegypti GN=AAEL006823 PE=4 SV=1
CCG005797.1	342	123	66	15.4540274	6.746334114	-1.195807182	Down	2.39E-08	4.71E-08	-
CCG000438.1	696	124	66	7.65552743	3.315009004	-1.207488987	Down	1.63E-08	3.22E-08	AAEL010298-PA OS=Aedes aegypti GN=AAEL010298 PE=4 SV=1
CCG017919.1	1131	128	66	4.86306457	2.040005541	-1.253292677	Down	3.43E-09	6.95E-09	AAEL013864-PA OS=Aedes aegypti GN=AAEL013864 PE=4 SV=1
CCG010043.1	951	128	66	5.78351843	2.426126464	-1.253292677	Down	3.43E-09	6.95E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017097 PE=4 SV=1
CCG016133.1	1641	129	66	3.37787676	1.406000163	-1.264519932	Down	2.31E-09	4.71E-09	AAEL006804-PA OS=Aedes aegypti GN=CYP9J18 PE=3 SV=2
CCG006997.1	377	148	66	16.8687552	6.120016623	-1.462746042	Down	8.02E-13	2.01E-12	AAEL014960-PA (Fragment) OS=Aedes aegypti GN=AAEL014960 PE=4 SV=1
CCG021014.1	2340	160	66	2.93810151	0.986002678	-1.575220772	Down	3.58E-15	1.29E-14	AAEL002031-PA OS=Aedes aegypti GN=CYP12F7 PE=3 SV=2
CCG021591.1	1164	163	66	6.01723947	1.982170332	-1.602020831	Down	8.88E-16	3.24E-15	AAEL008165-PA OS=Aedes aegypti GN=AAEL008165 PE=4 SV=1
CCG003838.2	3168	305	66	4.13692205	0.728297433	-2.505958109	Down	1.11E-49	5.15E-49	AAEL001687-PA OS=Aedes aegypti GN=AAEL001687 PE=4 SV=1

CCG008809.1	4338	930	66	9.21204546	0.531868665	-4.114379583	Down	2.31E-237	1.42E-236	AAEL007226-PA OS=Aedes aegypti GN=AAEL007226 PE=4 SV=1
CCG006003.1	945	1	67	0.04547062	2.478523327	5.768402394	Up	0.000587	0.00091831	-
CCG017929.1	693	1	67	0.06200539	3.379804536	5.768402394	Up	0.000587	0.00091791	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008409 PE=4 SV=1
CCG017509.1	1284	8	67	0.2677242	1.824146841	2.768402394	Up	2.19E-06	3.96E-06	AAEL000774-PA OS=Aedes aegypti GN=AAEL000774 PE=4 SV=1
CCG023660.1	2076	9	67	0.18628498	1.128229549	2.598477393	Up	1.10E-06	2.01E-06	AAEL008045-PA OS=Aedes aegypti GN=AAEL008045 PE=4 SV=1
CCG005459.1	918	10	67	0.4680799	2.551421072	2.446474299	Up	5.45E-07	1.01E-06	AAEL005327-PA OS=Aedes aegypti GN=AAEL005327 PE=4 SV=1
CCG021927.1	450	10	67	0.95488299	5.204898986	2.446474299	Up	5.45E-07	1.01E-06	AAEL000284-PA OS=Aedes aegypti GN=AAEL000284 PE=4 SV=1
CCG004739.1	1392	10	67	0.30869062	1.682618207	2.446474299	Up	5.45E-07	1.01E-06	AAEL000683-PA OS=Aedes aegypti GN=AAEL000683 PE=4 SV=1
CCG028188.1	780	12	67	0.66107284	3.002826338	2.183439894	Up	1.57E-07	2.98E-07	-
CCG027184.1	396	14	67	1.51913203	5.914657939	1.961047472	Up	2.35E-07	4.44E-07	AAEL008932-PA OS=Aedes aegypti GN=AAEL008932 PE=3 SV=1
CCG025672.1	736	15	67	0.87574187	3.18234313	1.861511799	Up	5.33E-07	9.95E-07	-
CCG014714.1	645	16	67	1.0659159	3.631324874	1.768402394	Up	1.26E-06	2.30E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ019728 PE=4 SV=1
CCG009824.1	1152	17	67	0.63410199	2.033163666	1.680939553	Up	2.87E-06	5.17E-06	-
CCG011829.1	2030	25	67	0.52918392	1.153795342	1.124546205	Up	0.0005478	0.00085857	AAEL010485-PA OS=Aedes aegypti GN=AAEL010485 PE=3 SV=1
CCG020996.1	921	122	67	5.69197353	2.543110254	-1.162334943	Down	5.54E-08	1.07E-07	Down syndrome cell adhesion molecule-like protein OS=Acromyrmex echinatio GN=G5I_10191 PE=4 SV=1
CCG014689.1	936	155	67	7.1157146	2.502355282	-1.507722011	Down	6.52E-14	2.14E-13	AAEL008646-PA (Fragment) OS=Aedes aegypti GN=AAEL008646 PE=4 SV=1

CCG020016.1	1805	206	67	4.90402511	1.297620246	-1.918098133	Down	9.63E-25	3.91E-24	Putative gag-pol protein OS=Drosophila ananassae PE=4 SV=1
CCG009167.1	1112	240	67	9.27404344	2.10629905	-2.138488201	Down	9.73E-33	4.16E-32	AAEL015038-PA OS=Aedes aegypti GN=AAEL015038 PE=4 SV=1
CCG025900.1	1563	314	67	8.63243549	1.498531378	-2.526218355	Down	1.44E-51	6.77E-51	AAEL009520-PA OS=Aedes aegypti GN=AAEL009520 PE=4 SV=1
CCG015684.1	864	323	67	16.063917	2.710884889	-2.56698796	Down	5.83E-54	2.76E-53	AAEL000689-PA OS=Aedes aegypti GN=AAEL000689 PE=3 SV=1
CCG005194.1	5469	323	67	2.53779928	0.428269253	-2.56698796	Down	5.83E-54	2.76E-53	AAEL012496-PA OS=Aedes aegypti GN=AAEL012496 PE=4 SV=1
CCG023321.3	1890	374	67	8.50300568	1.239261663	-2.778492066	Down	7.66E-68	3.78E-67	AAEL008342-PA OS=Aedes aegypti GN=AAEL008342 PE=4 SV=1
CCG019899.1	801	405	67	21.7262703	2.924100554	-2.893375703	Down	1.65E-76	8.29E-76	Hydroxypyruvate isomerase OS=Aedes aegypti GN=AAEL002165 PE=3 SV=1
CCG025108.1	2646	510	67	8.28214839	0.885186902	-3.225951043	Down	7.68E-107	4.10E-106	AAEL012077-PA (Fragment) OS=Aedes aegypti GN=AAEL012077 PE=3 SV=1
CCG014698.1	1041	718	67	29.6371464	2.249956334	-3.71943764	Down	7.45E-170	4.32E-169	AAEL010344-PA OS=Aedes aegypti GN=AAEL010344 PE=4 SV=1
CCG003567.2	933	764	67	35.1863636	2.51040144	-3.809026434	Down	3.82E-184	2.25E-183	AAEL001293-PA OS=Aedes aegypti GN=AAEL001293 PE=4 SV=1
CCG016766.1	810	1105	67	58.6192059	2.891610548	-4.34142826	Down	1.59E-292	9.97E-292	AAEL013542-PA OS=Aedes aegypti GN=AAEL013542 PE=4 SV=1
CCG020270.1	459	3106	67	290.771232	5.102842143	-5.83243972	Down	0	0	AAEL002185-PA OS=Aedes aegypti GN=AAEL002185 PE=4 SV=1
CCG007086.1	631	0	68	0.001	3.767294486	11.8793131	Up	4.44E-16	1.65E-15	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG014169.1	855	4	68	0.201028	2.780307392	3.789776045	Up	2.95E-05	5.04E-05	-
CCG007951.1	1578	12	68	0.32676604	1.506440317	2.204813544	Up	1.48E-07	2.81E-07	AAEL005945-PA (Fragment) OS=Aedes aegypti GN=GPRDOP4 PE=3 SV=2

CCG010495.1	598	12	68	0.86226892	3.975188663	2.204813544	Up	1.48E-07	2.81E-07	AAEL003105-PA (Fragment) OS=Aedes aegypti GN=AAEL003105 PE=4 SV=1
CCG028427.1	2019	20	68	0.42565364	1.177396147	1.46784795	Up	1.90E-05	3.27E-05	Proacrosin OS=Culex quinquefasciatus GN=CpipJ_CPIJ012481 PE=3 SV=1
CCG006280.1	669	20	68	1.28459595	3.553307654	1.46784795	Up	1.90E-05	3.27E-05	AAEL000949-PA (Fragment) OS=Aedes aegypti GN=AAEL000949 PE=4 SV=1
CCG000404.1	942	20	68	0.91230859	2.52352741	1.46784795	Up	1.90E-05	3.27E-05	AAEL004392-PA OS=Aedes aegypti GN=IMP PE=4 SV=1
CCG006844.1	501	22	68	1.88689453	4.744835969	1.330344426	Up	7.13E-05	0.00011825	Uncharacterized protein OS=Anopheles darlingi GN=AND_12274 PE=4 SV=1
CCG001057.1	984	117	68	5.10920625	2.415815875	-1.080588675	Down	5.30E-07	9.89E-07	Phosphatidate cytidyltransferase OS=Anopheles gambiae GN=AGAP007175 PE=3 SV=4
CCG005269.1	696	168	68	10.3720049	3.415463823	-1.602541378	Down	3.18E-16	1.19E-15	AAEL000254-PA OS=Aedes aegypti GN=AAEL000254 PE=4 SV=1
CCG003270.1	759	185	68	10.473519	3.131966825	-1.741605415	Down	9.05E-20	3.52E-19	GJ11154 OS=Drosophila virilis GN=GJ11154 PE=4 SV=1
CCG015674.1	1560	207	68	5.70175325	1.523822321	-1.903710912	Down	1.28E-24	5.20E-24	AAEL001442-PA OS=Aedes aegypti GN=AAEL001442 PE=4 SV=1
CCG015379.1	1657	234	68	6.06814599	1.434618479	-2.080588675	Down	6.72E-31	2.85E-30	AAEL013421-PA OS=Aedes aegypti GN=AAEL013421 PE=3 SV=1
CCG025946.1	1227	342	68	11.9768942	1.937378012	-2.62807647	Down	1.51E-58	7.24E-58	AAEL003560-PA OS=Aedes aegypti GN=AAEL003560 PE=3 SV=1
CCG006316.3	1080	407	68	16.1932241	2.201076686	-2.879108939	Down	1.74E-76	8.74E-76	AAEL006344-PA OS=Aedes aegypti GN=AAEL006344 PE=4 SV=1
CCG028497.1	918	1	69	0.04680799	2.627582895	5.810837661	Up	0.000587	0.00091677	AAEL012339-PA OS=Aedes aegypti GN=AAEL012339 PE=4 SV=1
CCG023453.1	628	4	69	0.27369258	3.840957161	3.810837661	Up	2.95E-05	5.04E-05	-
CCG003233.1	1033	5	69	0.20798516	2.335063986	3.488909566	Up	1.59E-05	2.76E-05	-
CCG004042.1	363	11	69	1.30211317	6.644961701	2.351406042	Up	2.71E-07	5.09E-07	AAEL006460-PA OS=Aedes aegypti GN=AAEL006460 PE=4 SV=1

CCG000391.1	8652	23	69	0.11422837	0.278793469	1.287275705	Up	9.55E-05	0.00015595	AAEL010367-PA OS=Aedes aegypti GN=AAEL010367 PE=4 SV=1
CCG026955.1	1404	25	69	0.7651306	1.71803497	1.166981471	Up	0.000302	0.00048041	AGAP005300-PA OS=Anopheles gambiae GN=AGAP005300 PE=2 SV=1
CCG020909.1	627	144	69	9.86864718	3.84708309	-1.359087341	Down	2.42E-11	5.28E-11	GTP-binding protein yptv3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004450 PE=3 SV=1
CCG010944.1	2955	161	69	2.34115982	0.816284635	-1.520079218	Down	1.49E-14	5.29E-14	AAEL012294-PA OS=Aedes aegypti GN=AAEL012294 PE=3 SV=1
CCG025963.1	750	333	69	19.0785622	3.216161463	-2.568540706	Down	1.29E-55	6.12E-55	AAEL003803-PA OS=Aedes aegypti GN=AAEL003803 PE=4 SV=1
CCG021804.1	1264	368	69	12.5101759	1.908323653	-2.712724295	Down	4.36E-65	2.13E-64	AAEL000249-PA OS=Aedes aegypti GN=AAEL000249 PE=3 SV=1
CCG000619.1	942	423	69	19.2953267	2.560638108	-2.913676193	Down	2.06E-80	1.05E-79	AAEL000723-PA OS=Aedes aegypti GN=AAEL006723 PE=4 SV=1
CCG013070.1	2055	3	70	0.06272954	1.190792883	4.24663372	Up	5.28E-05	8.86E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018868 PE=4 SV=1
CCG010160.1	402	6	70	0.64133932	6.087262125	3.24663372	Up	8.38E-06	1.48E-05	AAEL008675-PA OS=Aedes aegypti GN=AAEL008675 PE=4 SV=1
CCG011412.1	945	13	70	0.59111804	2.589501983	2.131156503	Up	8.44E-08	1.62E-07	AAEL009634-PA OS=Aedes aegypti GN=AAEL009634 PE=3 SV=1
CCG017616.1	1116	16	70	0.61605354	2.192723453	1.831596221	Up	3.90E-07	7.30E-07	AAEL015070-PA OS=Aedes aegypti GN=ALP PE=2 SV=1
CCG013901.1	665	16	70	1.03385828	3.679818608	1.831596221	Up	3.90E-07	7.30E-07	AAEL004635-PA OS=Aedes aegypti GN=AAEL004635 PE=4 SV=1
CCG023504.1	725	17	70	1.00756619	3.375281895	1.74413338	Up	9.12E-07	1.68E-06	AAEL005483-PA OS=Aedes aegypti GN=AAEL005483 PE=4 SV=1
CCG003884.1	1698	17	70	0.43020347	1.441153931	1.74413338	Up	9.12E-07	1.68E-06	AAEL012956-PA OS=Aedes aegypti GN=AAEL012956 PE=3 SV=1
CCG027671.1	2481	18	70	0.3117514	0.986327841	1.661671219	Up	2.06E-06	3.75E-06	AAEL010614-PA OS=Aedes aegypti GN=AAEL010614 PE=4 SV=1
CCG016975.1	1251	18	70	0.61826956	1.95609862	1.661671219	Up	2.06E-06	3.75E-06	AAEL011210-PA OS=Aedes aegypti GN=AAEL011210 PE=4 SV=1

CCG014818.1	282	22	70	3.3522488	8.677586433	1.372164602	Up	3.68E-05	6.21E-05	AAEL003171-PA OS=Aedes aegypti GN=AAEL003171 PE=4 SV=1
CCG004717.1	778	23	70	1.27031349	3.145346239	1.308034265	Up	6.91E-05	0.00011472	-
CCG009937.1	1206	26	70	0.92637902	2.029087375	1.131156503	Up	0.0003837	0.00060713	AAEL004556-PA OS=Aedes aegypti GN=AAEL004556 PE=4 SV=1
CCG016813.1	1033	128	70	5.32442016	2.368905493	-1.168403779	Down	2.29E-08	4.50E-08	AAEL008651-PA OS=Aedes aegypti GN=AAEL008651 PE=4 SV=1
CCG010130.2	879	130	70	6.35502332	2.783935579	-1.190771592	Down	1.07E-08	2.14E-08	AAEL003929-PA OS=Aedes aegypti GN=AAEL003929 PE=4 SV=1
CCG011769.1	1839	138	70	3.22448253	1.330657626	-1.276928236	Down	4.74E-10	9.87E-10	AAEL011809-PA OS=Aedes aegypti GN=AAEL011809 PE=3 SV=1
CCG001674.1	1440	169	70	5.0429758	1.699360676	-1.569283216	Down	7.30E-16	2.67E-15	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018073 PE=4 SV=1
CCG023486.1	1234	257	70	8.94912625	1.983046494	-2.174028328	Down	1.18E-35	5.14E-35	Putative uncharacterized protein GLEAN_15470 OS=Tribolium castaneum GN=GLEAN_15470 PE=4 SV=1
CCG008125.1	3108	449	70	6.20766115	0.787348576	-2.978975414	Down	3.36E-87	1.74E-86	AAEL004214-PA (Fragment) OS=Aedes aegypti GN=AAEL004214 PE=4 SV=1
CCG021146.1	477	21879	70	1970.93254	5.130145438	-8.585662959	Down	0	0	AAEL011741-PB OS=Aedes aegypti GN=GSTS1 PE=3 SV=2
CCG028184.1	525	0	71	0.001	4.727690763	12.20691996	Up	4.44E-16	1.64E-15	Putative uncharacterized protein OS=Aedes triseriatus PE=2 SV=1
CCG009757.1	396	2	71	0.21701886	6.267771846	4.852060323	Up	9.36E-05	0.0001539	AAEL006197-PA OS=Aedes aegypti GN=AAEL006197 PE=4 SV=1
CCG014142.1	291	5	71	0.73831159	8.529339006	3.530132228	Up	1.59E-05	2.76E-05	AGAP000639-PA OS=Anopheles gambiae GN=AgaP_AGAP000639 PE=3 SV=5
CCG018287.1	381	6	71	0.67668873	6.514534517	3.267097823	Up	8.38E-06	1.48E-05	Histone H2A OS=Aedes aegypti GN=AAEL003669 PE=3 SV=1
CCG014915.1	1047	7	71	0.28728571	2.370618578	3.044705401	Up	4.31E-06	7.69E-06	AAEL006329-PA OS=Aedes aegypti GN=AAEL006329 PE=4 SV=1
CCG005532.1	814	8	71	0.42230697	3.049186303	2.852060323	Up	2.19E-06	3.96E-06	-

CCG015202.1	732	8	71	0.46961459	3.390761818	2.852060323	Up	2.19E-06	3.96E-06	AAEL010866-PA OS=Aedes aegypti GN=AAEL010866 PE=3 SV=1
CCG001096.1	603	13	71	0.92637902	4.116148675	2.151620605	Up	7.70E-08	1.48E-07	Coactivator required for Sp1 transcriptional activation subunit 9 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007281 PE=4 SV=1
CCG026307.1	636	13	71	0.87831219	3.902574923	2.151620605	Up	7.70E-08	1.48E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003129 PE=4 SV=1
CCG018041.1	564	13	71	0.99043715	4.400775977	2.151620605	Up	7.70E-08	1.48E-07	AAEL012187-PA OS=Aedes aegypti GN=AAEL012187 PE=4 SV=1
CCG000499.1	1224	15	71	0.52658988	2.027808538	1.945169728	Up	1.17E-07	2.23E-07	Uncharacterized protein OS=Anopheles darlingi GN=AND_14401 PE=4 SV=1
CCG022645.1	525	15	71	1.2277067	4.727690763	1.945169728	Up	1.17E-07	2.23E-07	GSG8-like protein OS=Aedes aegypti PE=2 SV=1
CCG010639.1	225	15	71	2.86464897	11.03127845	1.945169728	Up	1.17E-07	2.23E-07	AAEL000719-PA OS=Aedes aegypti GN=AAEL000719 PE=4 SV=1
CCG023143.1	957	18	71	0.80820817	2.593560764	1.682135322	Up	1.42E-06	2.59E-06	AAEL006426-PA OS=Aedes aegypti GN=AAEL006426 PE=4 SV=1
CCG026943.1	575	23	71	1.71878938	4.316587219	1.328498367	Up	4.99E-05	8.38E-05	-
CCG019534.1	1539	24	71	0.67009333	1.612760007	1.267097823	Up	9.19E-05	0.00015144	AAEL010406-PA OS=Aedes aegypti GN=AAEL010406 PE=4 SV=1
CCG008380.1	1534	25	71	0.70028903	1.618016722	1.208204134	Up	0.0001642	0.0002649	AAEL003987-PA OS=Aedes aegypti GN=AAEL003987 PE=4 SV=1
CCG016019.1	1035	121	71	5.02351487	2.39810401	-1.066802914	Down	4.49E-07	8.39E-07	AAEL010041-PA OS=Aedes aegypti GN=AAEL010041 PE=4 SV=1
CCG018206.1	2169	123	71	2.4367346	1.14432349	-1.090454182	Down	2.21E-07	4.18E-07	AAEL014140-PA OS=Aedes aegypti GN=CLIPB24 PE=3 SV=1
CCG018782.1	1455	137	71	4.04594752	1.705867801	-1.24597176	Down	1.16E-09	2.38E-09	AAEL005243-PA OS=Aedes aegypti GN=AAEL005243 PE=4 SV=1
CCG011172.1	966	141	71	6.2719799	2.569397154	-1.287491029	Down	2.36E-10	4.98E-10	Annexin OS=Aedes aegypti GN=AAEL005412 PE=3 SV=1

CCG008388.1	919	147	71	6.87328726	2.700802667	-1.347612022	Down	2.04E-11	4.46E-11	Aminomethyltransferase OS=Aedes aegypti GN=AAEL010276 PE=3 SV=1
CCG017856.1	1308	169	71	5.55189996	1.897582302	-1.548819113	Down	1.38E-15	5.00E-15	AAEL012486-PA OS=Aedes aegypti GN=AAEL012486 PE=4 SV=1
CCG024068.1	522	180	71	14.8171499	4.7548614	-1.639792773	Down	8.24E-18	3.14E-17	AAEL008638-PA OS=Aedes aegypti GN=CYP49A1 PE=3 SV=1
CCG022587.1	2570	227	71	3.79538123	0.965773405	-1.974488164	Down	4.02E-28	1.68E-27	AAEL013884-PA OS=Aedes aegypti GN=AAEL013884 PE=4 SV=1
CCG009616.1	1032	276	71	11.4919058	2.405075243	-2.256464133	Down	6.28E-40	2.80E-39	AAEL009632-PA OS=Aedes aegypti GN=AAEL009632 PE=3 SV=1
CCG002848.1	727	327	71	19.3275147	3.414082051	-2.501086502	Down	4.98E-53	2.35E-52	AAEL003464-PA OS=Aedes aegypti GN=AAEL003464 PE=4 SV=1
CCG002683.1	1419	457	71	13.8387376	1.749145631	-2.983990032	Down	7.12E-89	3.69E-88	Uncharacterized protein OS=Anopheles darlingi GN=AND_03450 PE=4 SV=1
CCG009376.1	702	514	71	31.4621704	3.535666169	-3.153564226	Down	2.36E-105	1.26E-104	AAEL005218-PA OS=Aedes aegypti GN=AAEL005218 PE=4 SV=1
CCG023444.1	1587	542	71	14.6752339	1.563980876	-3.230088718	Down	1.42E-113	7.69E-113	AAEL002669-PA OS=Aedes aegypti GN=AAEL002669 PE=4 SV=1
CCG021005.1	1689	663	71	16.8673381	1.469530877	-3.520804737	Down	7.52E-150	4.27E-149	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003199 PE=4 SV=1
CCG017305.1	720	856	71	51.08624	3.447274515	-3.889406663	Down	1.55E-209	9.27E-209	AAEL006598-PA OS=Aedes aegypti GN=AAEL006598 PE=3 SV=1
CCG006375.1	1099	0	72	0.001	2.290260171	11.16129578	Up	4.44E-16	1.63E-15	Peritrophin OS=Aedes aegypti PE=2 SV=1
CCG009873.1	603	6	72	0.42755955	4.1741226	3.287275705	Up	8.38E-06	1.48E-05	AAEL000739-PA OS=Aedes aegypti GN=AAEL000739 PE=4 SV=1
CCG011452.1	1448	11	72	0.32642754	1.738256856	2.412806587	Up	2.68E-07	5.04E-07	AAEL012478-PA OS=Aedes aegypti GN=AAEL012478 PE=4 SV=1
CCG025406.1	2082	12	72	0.24766418	1.208931762	2.287275705	Up	1.33E-07	2.54E-07	Starlet OS=Culex quinquefasciatus GN=CpipJ_CPIJ003276 PE=2 SV=1
CCG014251.1	1863	15	72	0.3459721	1.351044513	1.96534761	Up	8.21E-08	1.58E-07	AAEL008523-PA OS=Aedes aegypti GN=AAEL008523 PE=4 SV=1

CCG004932.1	1071	15	72	0.60181701	2.350136254	1.96534761	Up	8.21E-08	1.58E-07	AAEL003007-PA OS=Aedes aegypti GN=AAEL003007 PE=4 SV=1
CCG004128.1	314	25	72	3.42115721	8.015910597	1.228382015	Up	0.0001205	0.00019594	LINE-1 retrotransposon-like element, putative (Fragment) OS=Aspergillus flavus (strain ATCC 200026 / FGSC A1120 / NRRL 3357 / JCM 12722 / SRRC 167) GN=AFLA_063980 PE=4 SV=1
CCG025572.1	1134	28	72	1.0609811	2.219573128	1.064883283	Up	0.0006002	0.00093387	AAEL001373-PA OS=Aedes aegypti GN=AAEL001373 PE=3 SV=1
CCG005679.1	597	141	72	10.1486308	4.216073581	-1.267313147	Down	3.90E-10	8.15E-10	AAEL001355-PA OS=Aedes aegypti GN=AAEL001355 PE=4 SV=1
CCG000602.1	1086	159	72	6.29114899	2.317675808	-1.44064475	Down	2.17E-13	6.21E-13	AAEL002010-PA OS=Aedes aegypti GN=AAEL002010 PE=3 SV=1
CCG012361.1	586	265	72	19.4317059	4.295214894	-2.177610344	Down	8.68E-37	3.80E-36	-
CCG009347.2	234	556	72	102.099028	10.75639285	-3.246702867	Down	5.00E-117	2.72E-116	AAEL002074-PA OS=Aedes aegypti GN=AAEL002074 PE=4 SV=1
CCG008675.1	1353	804	72	25.5341217	1.860307411	-3.778813486	Down	1.87E-192	1.11E-191	AAEL007575-PA OS=Aedes aegypti GN=AAEL007575 PE=4 SV=1
CCG002051.1	870	1113	72	54.971626	2.893098767	-4.247999672	Down	2.95E-290	1.85E-289	AAEL002089-PA OS=Aedes aegypti GN=AAEL002089 PE=4 SV=1
CCG014491.1	975	2	73	0.08814305	2.617388928	4.892137763	Up	9.36E-05	0.00015401	AAEL000529-PA OS=Aedes aegypti GN=AAEL000529 PE=4 SV=1
CCG010858.1	567	4	73	0.30313746	4.500801066	3.892137763	Up	2.95E-05	5.03E-05	Uncharacterized protein OS=Anopheles darlingi GN=AND_00155 PE=4 SV=1
CCG027027.1	186	5	73	1.15510039	13.72018389	3.570209668	Up	1.59E-05	2.76E-05	-
CCG027866.1	1539	12	73	0.33504666	1.658189866	2.307175262	Up	1.32E-07	2.52E-07	AAEL012391-PA OS=Aedes aegypti GN=AAEL012391 PE=4 SV=1
CCG018271.1	2094	12	73	0.2462449	1.218698283	2.307175262	Up	1.32E-07	2.52E-07	Monocarboxylate transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ006419 PE=4 SV=1
CCG014565.1	621	15	73	1.03791629	4.10942706	1.985247167	Up	5.93E-08	1.15E-07	AAEL011990-PA (Fragment) OS=Aedes aegypti GN=AAEL011990 PE=4 SV=1

CCG004123.1	2382	18	73	0.32470832	1.071349372	1.722212761	Up	6.61E-07	1.23E-06	AAEL007835-PA OS=Aedes aegypti GN=AAEL007835 PE=4 SV=1
CCG022233.1	504	23	73	1.96092043	5.063401199	1.368575807	Up	2.58E-05	4.41E-05	AAEL014499-PA OS=Aedes aegypti GN=AAEL014499 PE=4 SV=1
CCG002389.1	1908	27	73	0.60806228	1.337502204	1.137250261	Up	0.000269	0.00042917	AAEL012036-PA OS=Aedes aegypti GN=AAEL012036 PE=4 SV=1
CCG002634.1	927	27	73	1.25154567	2.752917157	1.137250261	Up	0.000269	0.0004292	AAEL010866-PA OS=Aedes aegypti GN=AAEL010866 PE=3 SV=1
CCG004552.1	828	127	73	6.59076847	3.082070295	-1.096546924	Down	1.23E-07	2.36E-07	AAEL001882-PA OS=Aedes aegypti GN=AAEL001882 PE=4 SV=1
CCG024998.1	1836	129	73	3.01911534	1.38995327	-1.119089493	Down	6.00E-08	1.16E-07	AAEL003653-PA OS=Aedes aegypti GN=SRPN12 PE=4 SV=2
CCG007684.1	2122	159	73	3.21969265	1.202617438	-1.420745193	Down	3.84E-13	1.03E-12	AAEL001515-PA OS=Aedes aegypti GN=AAEL001515 PE=4 SV=1
CCG024069.1	1026	173	73	7.2453841	2.4872848	-1.542490465	Down	7.80E-16	2.84E-15	AAEL008638-PA OS=Aedes aegypti GN=CYP49A1 PE=3 SV=1
CCG004335.1	1785	180	73	4.33308248	1.429666221	-1.599715334	Down	3.10E-17	1.18E-16	AAEL011716-PA OS=Aedes aegypti GN=AAEL011716 PE=4 SV=1
CCG005330.1	1875	221	73	5.06469939	1.361042242	-1.895764797	Down	4.89E-26	2.00E-25	AAEL013271-PA OS=Aedes aegypti GN=AAEL013271 PE=4 SV=1
CCG022286.1	1515	230	73	6.52345806	1.684458221	-1.953352288	Down	4.36E-28	1.81E-27	AGAP004228-PA OS=Anopheles gambiae GN=AgaP_AGAP004228 PE=4 SV=4
CCG018923.1	2807	270	73	4.13317718	0.909139367	-2.184677834	Down	1.34E-37	5.91E-37	AAEL008483-PA OS=Aedes aegypti GN=AAEL008483 PE=4 SV=1
CCG026384.1	3336	289	73	3.72249799	0.764974282	-2.28278792	Down	2.58E-42	1.16E-41	AAEL002632-PA OS=Aedes aegypti GN=AAEL002632 PE=3 SV=1
CCG016304.1	225	453	73	86.512399	11.34201869	-2.931229477	Down	1.59E-86	8.19E-86	AAEL006000-PA OS=Aedes aegypti GN=AAEL006000 PE=4 SV=1
CCG028235.1	567	2	74	0.15156873	4.562455875	4.911766569	Up	9.36E-05	0.00015368	AAEL007059-PA OS=Aedes aegypti GN=PGRPS5_b PE=4 SV=1
CCG027029.1	676	3	74	0.19069409	3.826793611	4.326804069	Up	5.28E-05	8.85E-05	-
CCG014770.1	252	5	74	0.8525741	10.26552572	3.589838475	Up	1.59E-05	2.76E-05	Transmembrane protein 42 OS=Acromyrmex echinatio GN=G5I_06843 PE=4 SV=1

CCG017578.1	912	6	74	0.28269562	2.836526843	3.326804069	Up	8.38E-06	1.48E-05	AAEL005379-PA OS=Aedes aegypti GN=AAEL005379 PE=4 SV=1
CCG010878.1	2823	17	74	0.25876213	0.91636999	1.824303728	Up	1.93E-07	3.65E-07	AAEL011292-PA OS=Aedes aegypti GN=AAEL011292 PE=4 SV=1
CCG027075.1	1110	17	74	0.65809503	2.330551785	1.824303728	Up	1.93E-07	3.65E-07	AAEL010525-PA OS=Aedes aegypti GN=AAEL010525 PE=4 SV=1
CCG027543.1	714	18	74	1.08327062	3.623126724	1.741841568	Up	4.50E-07	8.41E-07	-
CCG013820.1	957	19	74	0.85310863	2.703147838	1.663839056	Up	1.02E-06	1.88E-06	Uncharacterized protein OS=Anopheles darlingi GN=AND_14401 PE=4 SV=1
CCG015370.1	1412	21	74	0.63906829	1.832090992	1.519449147	Up	4.65E-06	8.28E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011805 PE=4 SV=1
CCG000663.1	1173	22	74	0.80591148	2.205381484	1.452334951	Up	9.42E-06	1.65E-05	AAEL011190-PA OS=Aedes aegypti GN=AAEL011190 PE=4 SV=1
CCG014554.1	992	24	74	1.03959035	2.607774679	1.326804069	Up	3.49E-05	5.91E-05	AAEL006197-PA OS=Aedes aegypti GN=AAEL006197 PE=4 SV=1
CCG022830.1	804	25	74	1.33612359	3.217552837	1.26791038	Up	6.43E-05	0.00010679	AAEL004953-PA OS=Aedes aegypti GN=AAEL004953 PE=4 SV=1
CCG002332.1	2016	25	74	0.53285881	1.283190715	1.26791038	Up	6.43E-05	0.0001068	AAEL000107-PA OS=Aedes aegypti GN=AAEL000107 PE=4 SV=1
CCG017845.1	327	28	74	3.67936565	7.911047343	1.104411647	Up	0.0003387	0.00053734	AAEL011433-PA OS=Aedes aegypti GN=AAEL011433 PE=4 SV=1
CCG018649.1	4364	28	74	0.27569949	0.592784712	1.104411647	Up	0.0003387	0.00053737	Reverse transcriptase homolog protein OS=Chironomus thummi GN=reverse transcriptase homolog PE=4 SV=1
CCG003623.1	723	121	74	7.19133871	3.578025562	-1.007096668	Down	1.49E-06	2.72E-06	AAEL012538-PA OS=Aedes aegypti GN=AAEL012538 PE=4 SV=1
CCG022151.1	559	133	74	10.2235683	4.627750414	-1.143515866	Down	2.15E-08	4.24E-08	AAEL002524-PA OS=Aedes aegypti GN=CTL24 PE=4 SV=1
CCG020127.1	1824	149	74	3.51013731	1.418263422	-1.307401951	Down	4.34E-11	9.35E-11	AAEL004491-PA OS=Aedes aegypti GN=AAEL004491 PE=4 SV=1
CCG021788.2	249	166	74	28.6464897	10.38920675	-1.463272862	Down	3.27E-14	1.13E-13	AAEL006602-PA OS=Aedes aegypti GN=AAEL006602 PE=4 SV=1

CCG012218.1	951	176	74	7.95233785	2.720202399	-1.547665049	Down	3.72E-16	1.39E-15	AAEL011460-PA OS=Aedes aegypti GN=AAEL011460 PE=4 SV=1
CCG019444.1	441	191	74	18.6104746	5.866014696	-1.665662259	Down	3.35E-19	1.30E-18	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005492 PE=4 SV=1
CCG010766.1	315	214	74	29.1921372	8.212420575	-1.829700417	Down	3.91E-24	1.58E-23	AAEL007194-PA OS=Aedes aegypti GN=AAEL007194 PE=4 SV=1
CCG020408.1	1788	219	74	5.26307152	1.446819061	-1.86302049	Down	3.03E-25	1.24E-24	AAEL007793-PA OS=Aedes aegypti GN=AAEL007793 PE=4 SV=1
CCG013931.2	1353	261	74	8.28906189	1.911982617	-2.116139427	Down	5.25E-35	2.27E-34	AAEL000500-PA OS=Aedes aegypti GN=AAEL000500 PE=4 SV=1
CCG021216.1	786	289	74	15.7993045	3.291237253	-2.263159113	Down	7.01E-42	3.15E-41	AGAP010534-PA (Fragment) OS=Anopheles gambiae GN=AGAP010534 PE=4 SV=3
CCG021525.1	1466	319	74	9.35016735	1.764606058	-2.405646044	Down	1.66E-49	7.71E-49	-
CCG024348.1	747	361	74	20.7658289	3.463068917	-2.584088457	Down	1.47E-60	7.10E-60	AAEL001306-PA OS=Aedes aegypti GN=AAEL001306 PE=4 SV=1
CCG005975.1	1430	555	74	16.6770648	1.809029707	-3.204577392	Down	2.21E-115	1.20E-114	AAEL001307-PA OS=Aedes aegypti GN=AAEL001307 PE=4 SV=1
CCG003309.1	1590	0	75	0.001	1.648975319	10.68735409	Up	4.44E-16	1.66E-15	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005832 PE=4 SV=1
CCG014487.1	354	2	75	0.24276686	7.40641457	4.931131894	Up	9.36E-05	0.00015373	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005832 PE=4 SV=1
CCG014411.1	810	2	75	0.10609811	3.236877479	4.931131894	Up	9.36E-05	0.00015348	Uncharacterized protein OS=Anopheles darlingi GN=AND_21799 PE=4 SV=1
CCG025928.1	705	8	75	0.48759983	3.718965614	2.931131894	Up	2.19E-06	3.96E-06	AAEL000666-PA OS=Aedes aegypti GN=AAEL000666 PE=4 SV=1
CCG020038.1	749	9	75	0.51632525	3.500495004	2.761206893	Up	1.09E-06	2.01E-06	-
CCG005020.1	873	13	75	0.63987005	3.003288382	2.230692176	Up	6.58E-08	1.27E-07	AAEL014040-PA OS=Aedes aegypti GN=AAEL014040 PE=4 SV=1

CCG000425.1	1336	15	75	0.48244463	1.962478112	2.024241299	Up	3.41E-08	6.66E-08	FA1556A OS=Culex quinquefasciatus GN=CpipJ_CPIJ004210 PE=4 SV=1
CCG008025.1	404	16	75	1.70177167	6.489779104	1.931131894	Up	5.79E-08	1.12E-07	AAEL003562-PA OS=Aedes aegypti GN=AAEL003562 PE=4 SV=1
CCG012506.1	729	17	75	1.00203771	3.596530532	1.843669053	Up	1.31E-07	2.49E-07	AAEL001543-PA OS=Aedes aegypti GN=AAEL001543 PE=4 SV=1
CCG019831.2	1380	20	75	0.62274978	1.899906346	1.609203799	Up	1.54E-06	2.80E-06	AAEL014938-PA OS=Aedes aegypti GN=AAEL014938 PE=3 SV=1
CCG015839.1	783	22	75	1.20732332	3.348493944	1.471700276	Up	6.65E-06	1.18E-05	Salivary antigen-5 related protein AG5-1 OS=Aedes albopictus PE=2 SV=1
CCG020374.1	1038	24	75	0.99351988	2.52588705	1.346169394	Up	2.52E-05	4.31E-05	AAEL006263-PA OS=Aedes aegypti GN=AAEL006263 PE=4 SV=1
CCG006269.1	1098	25	75	0.97836372	2.387860435	1.287275705	Up	4.67E-05	7.85E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014761 PE=4 SV=1
CCG014126.1	778	125	75	6.90387767	3.370013828	-1.03465239	Down	5.66E-07	1.05E-06	Gag-Pol polyprotein OS=Schistosoma japonicum PE=4 SV=1
CCG019578.1	1444	128	75	3.80895154	1.815699971	-1.068868106	Down	2.00E-07	3.78E-07	Pof-like protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012062 PE=4 SV=1
CCG012347.1	657	223	75	14.5848566	3.990670864	-1.869768006	Down	8.50E-26	3.48E-25	AAEL000457-PA OS=Aedes aegypti GN=AAEL000457 PE=4 SV=1
CCG005163.1	1551	289	75	8.00661077	1.690438915	-2.243793788	Down	1.89E-41	8.47E-41	AAEL001737-PA (Fragment) OS=Aedes aegypti GN=AAEL001737 PE=4 SV=1
CCG025788.1	1331	778	75	25.1167945	1.969850306	-3.672494451	Down	5.67E-182	3.33E-181	AAEL009480-PA OS=Aedes aegypti GN=AAEL009480 PE=4 SV=1
CCG023922.1	861	836	75	41.7220652	3.04514606	-3.776227238	Down	6.22E-200	3.71E-199	AAEL012650-PA OS=Aedes aegypti GN=AAEL012650 PE=4 SV=1
CCG009456.1	513	0	76	0.001	5.179003966	12.33845895	Up	4.44E-16	1.63E-15	-
CCG025027.1	588	0	76	0.001	4.518416726	12.14160162	Up	4.44E-16	1.63E-15	AAEL015271-PA OS=Aedes aegypti GN=AAEL015271 PE=3 SV=1

CCG008345.2	921	9	76	0.41989969	2.884722079	2.780315716	Up	1.09E-06	2.01E-06	AAEL002945-PA OS=Aedes aegypti GN=AAEL002945 PE=4 SV=2
CCG011462.1	2559	13	76	0.21829095	1.0382294	2.249800999	Up	6.49E-08	1.25E-07	AAEL010851-PA OS=Aedes aegypti GN=AAEL010851 PE=4 SV=1
CCG007491.1	822	13	76	0.67957001	3.23215211	2.249800999	Up	6.49E-08	1.25E-07	-
CCG025810.1	602	30	76	2.1413489	4.413337267	1.043350122	Up	0.0005216	0.00081859	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001634 PE=4 SV=1
CCG004834.1	2595	152	76	2.51691702	1.023826218	-1.297686796	Down	3.60E-11	7.81E-11	AAEL010111-PA OS=Aedes aegypti GN=AAEL010111 PE=4 SV=1
CCG025777.1	1467	168	76	4.9208694	1.811062737	-1.442076706	Down	4.38E-14	1.49E-13	AAEL002547-PA OS=Aedes aegypti GN=AAEL002547 PE=4 SV=1
CCG000758.1	3284	178	76	2.32905382	0.80902224	-1.525492714	Down	5.22E-16	1.91E-15	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012060 PE=4 SV=1
CCG002256.1	996	197	76	8.49903385	2.667499031	-1.671811102	Down	7.46E-20	2.91E-19	AAEL008108-PA OS=Aedes aegypti GN=AAEL008108 PE=4 SV=1
CCG017846.1	591	274	76	19.9216705	4.4954806	-2.147791366	Down	2.40E-37	1.06E-36	AAEL011439-PA OS=Aedes aegypti GN=AAEL011439 PE=4 SV=1
CCG011867.1	2247	297	76	5.67957774	1.182389424	-2.264078404	Down	5.14E-43	2.32E-42	AAEL011366-PA OS=Aedes aegypti GN=AAEL011366 PE=4 SV=1
CCG003934.1	1473	363	76	10.5892829	1.803685699	-2.553585021	Down	4.43E-60	2.14E-59	AAEL013039-PA OS=Aedes aegypti GN=AAEL013039 PE=4 SV=1
CCG003566.1	1353	689	76	21.881853	1.963657823	-3.478119455	Down	5.35E-154	3.05E-153	AAEL001307-PA OS=Aedes aegypti GN=AAEL001307 PE=4 SV=1
CCG012255.1	465	2913	76	269.184596	5.713610827	-5.558049269	Down	0	0	AAEL014419-PA OS=Aedes aegypti GN=AAEL014419 PE=4 SV=1
CCG007674.1	519	0	77	0.001	5.186488076	12.34054226	Up	4.44E-16	1.64E-15	-
CCG011919.1	7555	2	77	0.01137518	0.356292166	4.969099745	Up	9.36E-05	0.00015333	-
CCG008537.1	1333	4	77	0.12894144	2.01934532	3.969099745	Up	2.95E-05	5.02E-05	-
CCG021509.1	810	5	77	0.26524528	3.323194212	3.64717165	Up	1.59E-05	2.76E-05	Senné proteinase stubble OS=Culex quinquefasciatus GN=CpipJ_CPIJ009890 PE=3 SV=1
CCG005018.1	845	10	77	0.50851757	3.185547114	2.64717165	Up	5.43E-07	1.01E-06	-

CCG010769.1	342	11	77	1.38206749	7.870723133	2.509668126	Up	2.67E-07	5.03E-07	AAEL007543-PA OS=Aedes aegypti GN=AAEL007543 PE=4 SV=1
CCG028301.1	345	12	77	1.49459946	7.802282062	2.384137244	Up	1.31E-07	2.49E-07	AAEL008038-PA OS=Aedes aegypti GN=AAEL008038 PE=4 SV=1
CCG017886.1	247	12	77	2.08759844	10.89792434	2.384137244	Up	1.31E-07	2.49E-07	Ahr1 hydrocarbon receptor nuclear translocator OS=Daphnia magna GN=DmagArnt1 PE=2 SV=1
CCG002928.1	306	13	77	1.8255116	8.79669056	2.268660026	Up	6.44E-08	1.24E-07	AAEL000315-PA OS=Aedes aegypti GN=AAEL000315 PE=4 SV=1
CCG025541.1	2883	15	77	0.22356782	0.933675793	2.062209149	Up	2.31E-08	4.54E-08	Uncharacterized protein OS=Danio rerio GN=LOC100537316 PE=4 SV=1
CCG024169.1	428	16	77	1.60634522	6.28922269	1.969099745	Up	2.94E-08	5.77E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001644 PE=4 SV=1
CCG012037.1	270	17	77	2.70550181	9.969582635	1.881636903	Up	6.04E-08	1.17E-07	AAEL010350-PA OS=Aedes aegypti GN=AAEL010350 PE=4 SV=1
CCG001731.1	778	18	77	0.99415838	3.459880863	1.799174743	Up	1.41E-07	2.68E-07	-
CCG018241.1	1434	26	77	0.77908863	1.877118069	1.268660026	Up	4.50E-05	7.57E-05	AAEL001391-PA OS=Aedes aegypti GN=AAEL001391 PE=4 SV=1
CCG011916.1	1105	31	77	1.20548577	2.436006617	1.014903434	Up	0.0006389	0.00099237	AAEL010261-PA OS=Aedes aegypti GN=AAEL010261 PE=4 SV=1
CCG010387.1	1086	155	77	6.13288109	2.478625517	-1.307024661	Down	1.79E-11	3.93E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_22349 PE=4 SV=1
CCG004548.1	330	167	77	21.7452899	8.156931247	-1.414604548	Down	1.20E-13	3.68E-13	AAEL011718-PA OS=Aedes aegypti GN=AAEL011718 PE=4 SV=1
CCG009199.1	1525	178	77	5.01548378	1.765106434	-1.506633686	Down	9.63E-16	3.51E-15	AAEL010344-PA OS=Aedes aegypti GN=AAEL010344 PE=4 SV=1
CCG007746.1	1491	228	77	6.57082461	1.805357016	-1.86379027	Down	3.13E-26	1.29E-25	AAEL000044-PA OS=Aedes aegypti GN=CYP325Q1 PE=3 SV=1
CCG020135.1	480	278	77	24.886638	5.607890232	-2.149841328	Down	6.52E-38	2.88E-37	-
CCG006464.1	375	681	77	78.0330381	7.178099497	-3.442411243	Down	7.68E-151	4.36E-150	-
CCG008900.1	1278	1359	77	45.6931685	2.106249852	-4.439229996	Down	0	0	AAEL000797-PA OS=Aedes aegypti GN=AAEL000797 PE=4 SV=1

CCG024135.1	705	0	78	0.001	3.867724239	11.91726922	Up	4.44E-16	1.65E-15	AAEL006460-PA OS=Aedes aegypti GN=AAEL006460 PE=4 SV=1
CCG006932.1	885	0	78	0.001	3.081068461	11.58921502	Up	4.44E-16	1.64E-15	AAEL008197-PA OS=Aedes aegypti GN=AAEL008197 PE=3 SV=1
CCG005309.1	342	2	78	0.251285	7.972940316	4.987715423	Up	9.36E-05	0.00015356	-
CCG016796.1	4693	3	78	0.0274684	0.581023991	4.402752922	Up	5.28E-05	8.83E-05	-
CCG011572.1	886	4	78	0.19399429	3.077590957	3.987715423	Up	2.95E-05	5.03E-05	AAEL006220-PA OS=Aedes aegypti GN=AAEL006220 PE=4 SV=1
CCG025755.1	917	4	78	0.18743614	2.97355026	3.987715423	Up	2.95E-05	5.03E-05	AAEL011846-PA OS=Aedes aegypti GN=AAEL011846 PE=4 SV=1
CCG005785.1	1356	6	78	0.19013157	2.010874328	3.402752922	Up	8.38E-06	1.48E-05	Aryl hydrocarbon binding protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007210 PE=4 SV=1
CCG026450.1	518	11	78	0.91248471	5.263987622	2.528283804	Up	2.67E-07	5.03E-07	AAEL003190-PA OS=Aedes aegypti GN=AAEL003190 PE=4 SV=1
CCG017557.1	1287	11	78	0.36726269	2.118683441	2.528283804	Up	2.67E-07	5.03E-07	AAEL008914-PA OS=Aedes aegypti GN=AAEL008914 PE=4 SV=1
CCG001574.1	213	15	78	3.02603765	12.80162248	2.080824827	Up	2.02E-08	3.98E-08	AAEL001369-PA OS=Aedes aegypti GN=AAEL001369 PE=4 SV=1
CCG022636.1	5505	18	78	0.14050049	0.495321633	1.817790421	Up	9.54E-08	1.83E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015717 PE=4 SV=1
CCG028000.5	1929	18	78	0.40096175	1.41355396	1.817790421	Up	9.54E-08	1.83E-07	Heat shock 70 Ca OS=Aedes aegypti GN=Hsp70Ca PE=3 SV=1
CCG023149.1	2115	22	78	0.44696651	1.289241413	1.528283804	Up	2.31E-06	4.17E-06	AAEL003580-PA OS=Aedes aegypti GN=CYP4H28 PE=3 SV=2
CCG022784.1	1776	23	78	0.55647742	1.535329723	1.464153467	Up	4.69E-06	8.35E-06	Tel1 (Fragment) OS=Drosophila virilis GN=Tel1 PE=4 SV=1
CCG000053.1	483	133	78	11.8322458	5.645436	-1.067567013	Down	1.19E-07	2.28E-07	AAEL014888-PA OS=Aedes aegypti GN=AAEL014888 PE=4 SV=1
CCG010163.1	1176	134	78	4.89621126	2.318661214	-1.078373768	Down	8.37E-08	1.61E-07	AAEL008321-PA OS=Aedes aegypti GN=AAEL008321 PE=4 SV=1
CCG012096.1	1848	135	78	3.13902282	1.475511682	-1.089100174	Down	5.87E-08	1.14E-07	Transporter OS=Aedes aegypti GN=AAEL010106 PE=3 SV=1
CCG010414.1	1350	235	78	7.47991677	2.019811547	-1.888801524	Down	1.86E-27	7.71E-27	-

CCG003262.1	735	241	78	14.089396	3.709857943	-1.925173914	Down	8.16E-29	3.42E-28	AAEL001483-PA OS=Aedes aegypti GN=AAEL001483 PE=4 SV=1
CCG019363.1	1791	261	78	6.26192112	1.522471015	-2.040190574	Down	1.89E-33	8.11E-33	AAEL005200-PA (Fragment) OS=Aedes aegypti GN=AAEL005200 PE=4 SV=2
CCG008323.1	879	323	78	15.7897887	3.102099645	-2.347674932	Down	1.07E-48	4.98E-48	Synapsin OS=Culex quinquefasciatus GN=CpipJ_CPIJ003281 PE=4 SV=1
CCG018929.1	1314	393	78	12.8516786	2.075148849	-2.63067008	Down	4.25E-67	2.09E-66	Cytochrome P450 OS=Aedes aegypti PE=2 SV=1
CCG008194.1	873	970	78	47.7441496	3.123419918	-3.934125514	Down	2.56E-239	1.57E-238	AAEL004251-PA OS=Aedes aegypti GN=AAEL004251 PE=4 SV=1
CCG014484.1	7615	1037	78	5.85155808	0.358075586	-4.030484756	Down	1.81E-260	1.12E-259	BEL12_AG transposon polyprotein OS=Anopheles gambiae GN=BEL12-AG transposon PE=4 SV=1
CCG021210.1	573	0	79	0.001	4.819727513	12.23473587	Up	4.44E-16	1.64E-15	AAEL004817-PA OS=Aedes aegypti GN=AAEL004817 PE=4 SV=1
CCG012830.1	1298	1	79	0.03310457	2.127660913	6.006093952	Up	0.000587	0.00091558	AAEL003526-PA OS=Aedes aegypti GN=AAEL003526 PE=4 SV=1
CCG008625.1	822	3	79	0.15682385	3.359737062	4.421131451	Up	5.28E-05	8.85E-05	AAEL014554-PA OS=Aedes aegypti GN=AAEL014554 PE=4 SV=1
CCG007228.1	1452	5	79	0.14796741	1.901999907	3.684165857	Up	1.59E-05	2.76E-05	Putative pol-like protein OS=Danaus plexippus GN=KGM_11377 PE=4 SV=1
CCG003611.1	1658	6	79	0.15549964	1.665683875	3.421131451	Up	8.38E-06	1.48E-05	AAEL010790-PA OS=Aedes aegypti GN=AAEL010790 PE=4 SV=1
CCG021157.1	462	6	79	0.5580485	5.977713994	3.421131451	Up	8.38E-06	1.47E-05	-
CCG025890.1	321	17	79	2.27565573	8.603438832	1.918631111	Up	2.87E-08	5.64E-08	AAEL010637-PA OS=Aedes aegypti GN=AAEL010637 PE=4 SV=1
CCG018744.1	1332	23	79	0.74196989	2.07335125	1.482531996	Up	3.31E-06	5.94E-06	AAEL001787-PA OS=Aedes aegypti GN=AAEL001787 PE=4 SV=1
CCG005449.1	1098	25	79	0.97836372	2.515212992	1.362237762	Up	1.27E-05	2.21E-05	AAEL011071-PA OS=Aedes aegypti GN=AAEL011071 PE=4 SV=1

CCG014701.1	798	29	79	1.56155677	3.460781786	1.148112957	Up	0.0001323	0.00021489	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018760 PE=4 SV=1
CCG009772.1	501	32	79	2.74457387	5.512382964	1.006093952	Up	0.000593	0.00092322	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007337 PE=4 SV=1
CCG027636.1	369	135	79	15.7206346	7.484292317	-1.070721645	Down	8.91E-08	1.71E-07	AAEL011837-PA OS=Aedes aegypti GN=AAEL011837 PE=4 SV=1
CCG026143.1	5265	174	79	1.4200824	0.524540145	-1.436849544	Down	1.83E-14	6.43E-14	AAEL003307-PA (Fragment) OS=Aedes aegypti GN=AAEL003307 PE=4 SV=1
CCG023765.1	549	205	79	16.045165	5.030425984	-1.673386148	Down	1.27E-20	4.98E-20	-
CCG016612.1	2031	230	79	4.86609501	1.359775414	-1.839396099	Down	5.33E-26	2.18E-25	Beta-galactosidase OS=Aedes aegypti GN=AAEL004580 PE=3 SV=1
CCG014342.1	741	233	79	13.511401	3.726995769	-1.858092193	Down	1.15E-26	4.74E-26	AAEL005770-PA OS=Aedes aegypti GN=AAEL005770 PE=4 SV=1
CCG010111.1	2646	385	79	6.25221006	1.04372784	-2.582620684	Down	1.97E-64	9.62E-64	AAEL012077-PA (Fragment) OS=Aedes aegypti GN=AAEL012077 PE=3 SV=1
CCG021359.1	1122	1563	79	59.8589084	2.461411644	-4.604008111	Down	0	0	AAEL011771-PA OS=Aedes aegypti GN=AAEL011771 PE=4 SV=1
CCG014652.1	285	3090	79	465.882386	9.690189	-5.587297171	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ019849 PE=4 SV=1
CCG007675.1	648	0	80	0.001	4.315836639	12.07542454	Up	4.44E-16	1.65E-15	AAEL011890-PA OS=Aedes aegypti GN=AAEL011890 PE=4 SV=1
CCG018430.1	402	1	80	0.10688989	6.956870999	6.024241299	Up	0.000587	0.00091689	AAEL006551-PA OS=Aedes aegypti GN=AAEL006551 PE=4 SV=1
CCG018012.1	645	5	80	0.33309872	4.335910297	3.702313204	Up	1.59E-05	2.76E-05	AAEL001477-PA OS=Aedes aegypti GN=AAEL001477 PE=4 SV=1
CCG017373.1	606	10	80	0.70907153	4.614954029	2.702313204	Up	5.43E-07	1.01E-06	AAEL010423-PA OS=Aedes aegypti GN=AAEL010423 PE=4 SV=1
CCG019506.1	1771	11	80	0.26689276	1.579142937	2.56480968	Up	2.67E-07	5.03E-07	AAEL006034-PA OS=Aedes aegypti GN=AAEL006034 PE=4 SV=1

CCG013401.1	252	15	80	2.5577223	11.09786564	2.117350703	Up	1.71E-08	3.37E-08	AAEL004408-PA OS=Aedes aegypti GN=AAEL004408 PE=4 SV=1
CCG023351.2	627	16	80	1.09651635	4.460386191	2.024241299	Up	1.35E-08	2.68E-08	AAEL012899-PA OS=Aedes aegypti GN=AAEL012899 PE=4 SV=1
CCG011567.1	1839	18	80	0.42058468	1.520751572	1.854316297	Up	4.40E-08	8.56E-08	AAEL000050-PA OS=Aedes aegypti GN=AAEL006050 PE=4 SV=1
CCG013925.1	414	20	80	2.07583259	6.755222565	1.702313204	Up	2.36E-07	4.46E-07	AAEL000050-PA OS=Aedes aegypti GN=AAEL000503 PE=4 SV=1
CCG014388.1	1344	27	80	0.86323128	2.080849808	1.269353797	Up	3.15E-05	5.35E-05	AAEL000050-PA OS=Aedes aegypti GN=AAEL006556 PE=4 SV=1
CCG025518.1	1497	134	80	3.84632227	1.868177783	-1.041847892	Down	1.89E-07	3.59E-07	AAEL009892-PA OS=Aedes aegypti GN=AAEL009892 PE=4 SV=1
CCG017652.1	1443	149	80	4.43693032	1.938088802	-1.194927222	Down	8.23E-10	1.70E-09	AAEL014965-PA OS=Aedes aegypti GN=AAEL014965 PE=4 SV=1
CCG015644.1	858	218	80	10.9177181	3.259512986	-1.743943026	Down	4.61E-23	1.85E-22	AAEL010102-PA OS=Aedes aegypti GN=AAEL010102 PE=4 SV=1
CCG013923.1	402	240	80	25.6535729	6.956870999	-1.882649297	Down	6.87E-28	2.85E-27	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012173 PE=4 SV=1
CCG014186.1	507	0	81	0.001	5.585050135	12.44735452	Up	4.44E-16	1.65E-15	AAEL014453-PA OS=Aedes aegypti GN=AAEL014453 PE=4 SV=1
CCG003434.1	764	0	81	0.001	3.706309448	11.85576763	Up	4.44E-16	1.64E-15	-
CCG027414.1	885	0	81	0.001	3.199571094	11.64366281	Up	4.44E-16	1.63E-15	AAEL000713-PA OS=Aedes aegypti GN=AAEL000713 PE=4 SV=1
CCG019835.1	810	1	81	0.05304906	3.495827677	6.042163207	Up	0.000587	0.00091814	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014520 PE=3 SV=1
CCG014989.1	373	3	81	0.34560108	7.591475653	4.457200706	Up	5.28E-05	8.83E-05	AGAP007322-PA OS=Anopheles gambiae GN=AGAP007322 PE=4 SV=1
CCG026035.1	723	4	81	0.23773021	3.916487439	4.042163207	Up	2.95E-05	5.04E-05	AAEL000379-PA OS=Aedes aegypti GN=AAEL005379 PE=4 SV=1
CCG014496.1	300	5	81	0.71616224	9.438734729	3.720235112	Up	1.59E-05	2.76E-05	AAEL000331-PA OS=Aedes aegypti GN=AAEL006331 PE=4 SV=1

CCG023128.1	1515	7	81	0.19854003	1.869056382	3.234808285	Up	4.31E-06	7.70E-06	AAEL015347-PA OS=Aedes aegypti GN=AAEL015347 PE=4 SV=1
CCG026174.1	3306	8	81	0.10398	0.856509503	3.042163207	Up	2.19E-06	3.97E-06	Guanylate cyclase OS=Culex quinquefasciatus GN=CpipJ_CPIJ009261 PE=3 SV=1
CCG005457.1	1632	9	81	0.23696545	1.735061531	2.872238205	Up	1.09E-06	2.01E-06	AAEL005335-PA (Fragment) OS=Aedes aegypti GN=AAEL005335 PE=4 SV=1
CCG024564.1	864	13	81	0.64653536	3.277338447	2.341723489	Up	6.36E-08	1.23E-07	Tensin OS=Culex quinquefasciatus GN=CpipJ_CPIJ005796 PE=4 SV=1
CCG002921.1	399	14	81	1.50770999	7.096793029	2.234808285	Up	3.12E-08	6.10E-08	AAEL002461-PA OS=Aedes aegypti GN=AAEL002461 PE=4 SV=1
CCG015593.1	2220	24	81	0.46453767	1.275504693	1.457200706	Up	3.31E-06	5.94E-06	AAEL003726-PA OS=Aedes aegypti GN=AAEL003726 PE=4 SV=1
CCG016345.1	462	25	81	2.32520209	6.129048525	1.398307017	Up	6.50E-06	1.15E-05	AAEL006168-PA OS=Aedes aegypti GN=CLIPB42 PE=4 SV=1
CCG021585.1	654	30	81	1.97108874	4.32969483	1.135272611	Up	0.0001248	0.00020283	AAEL002836-PA (Fragment) OS=Aedes aegypti GN=AAEL002836 PE=4 SV=1
CCG002674.1	1445	32	81	0.9515789	1.959598906	1.042163207	Up	0.0003429	0.00054388	AAEL011199-PA OS=Aedes aegypti GN=AAEL011199 PE=4 SV=1
CCG017701.1	1812	142	81	3.36738538	1.562704425	-1.107583913	Down	1.72E-08	3.39E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007371 PE=4 SV=1
CCG027312.1	777	157	81	8.68243029	3.644299123	-1.252457542	Down	6.03E-11	1.30E-10	AAEL004126-PA OS=Aedes aegypti GN=AAEL004126 PE=4 SV=1
CCG001472.1	315	169	81	23.0536036	8.98927117	-1.35871623	Down	4.67E-13	1.22E-12	AGAP007867-PA (Fragment) OS=Anopheles gambiae GN=AGAP007867 PE=4 SV=3
CCG027424.1	855	182	81	9.14677392	3.311836747	-1.465631433	Down	1.79E-15	6.50E-15	CD109 antigen OS=Culex quinquefasciatus GN=CpipJ_CPIJ020143 PE=4 SV=1
CCG022403.1	843	184	81	9.37892191	3.35898033	-1.481398749	Down	7.42E-16	2.71E-15	AAEL013324-PA OS=Aedes aegypti GN=AAEL013324 PE=4 SV=1

CCG004543.1	549	187	81	14.6363213	5.15777854	-1.504731253	Down	1.96E-16	7.34E-16	AAEL000455-PA OS=Aedes aegypti GN=AAEL000455 PE=4 SV=1
CCG010263.1	1493	248	81	7.13763844	1.896597735	-1.912033104	Down	2.40E-29	1.01E-28	-
CCG000610.1	603	268	81	19.0976598	4.695887925	-2.023925984	Down	5.84E-34	2.52E-33	AAEL012400-PA OS=Aedes aegypti GN=AAEL012400 PE=4 SV=1
CCG024008.1	2094	282	81	5.78675509	1.352254259	-2.097388146	Down	2.81E-37	1.23E-36	AAEL000427-PA OS=Aedes aegypti GN=AAEL000427 PE=4 SV=1
CCG014356.1	1527	283	81	7.96361159	1.854368316	-2.102495036	Down	1.62E-37	7.12E-37	AAEL003966-PA OS=Aedes aegypti GN=AAEL003966 PE=4 SV=1
CCG015472.1	516	310	81	25.8151506	5.48763647	-2.233961199	Down	4.16E-44	1.89E-43	-
CCG012915.1	1260	328	81	11.1857722	2.247317793	-2.315388798	Down	1.28E-48	5.94E-48	AAEL005296-PA OS=Aedes aegypti GN=AAEL005296 PE=4 SV=1
CCG028536.2	2050	371	81	7.77647392	1.381278253	-2.49311217	Down	1.02E-59	4.89E-59	AAEL004544-PA OS=Aedes aegypti GN=AAEL004544 PE=4 SV=1
CCG009022.2	1911	512	81	11.512561	1.481747995	-2.957836793	Down	1.78E-98	9.42E-98	Transporter OS=Aedes aegypti GN=NAT8 PE=2 SV=1
CCG013258.1	1611	747	81	19.9245138	1.75767872	-3.502801226	Down	7.47E-168	4.33E-167	AAEL001233-PA OS=Aedes aegypti GN=CLYPE9 PE=3 SV=2
CCG013745.1	7683	928	81	5.19014886	0.368556608	-3.815817788	Down	1.32E-223	8.01E-223	AAEL000307-PA (Fragment) OS=Aedes aegypti GN=AAEL000307 PE=4 SV=1
CCG010923.1	852	3	82	0.15130188	3.364528985	4.474902708	Up	5.28E-05	8.85E-05	Osims OS=Culex quinquefasciatus GN=CpipJ_CPIJ008140 PE=4 SV=1
CCG020174.1	387	4	82	0.44413162	7.407180091	4.059865208	Up	2.95E-05	5.02E-05	-
CCG027808.1	1050	6	82	0.24554134	2.730074948	3.474902708	Up	8.38E-06	1.47E-05	AAEL000411-PA OS=Aedes aegypti GN=AAEL000411 PE=4 SV=1
CCG009228.1	501	10	82	0.85767933	5.721713963	2.737937114	Up	5.43E-07	1.01E-06	AAEL003000-PA OS=Aedes aegypti GN=AAEL003000 PE=2 SV=1
CCG020604.1	606	11	82	0.77997868	4.73032788	2.60043359	Up	2.67E-07	5.03E-07	AAEL002151-PA OS=Aedes aegypti GN=AAEL002151 PE=4 SV=1
CCG011444.1	558	18	82	1.38612047	5.137237805	1.889940207	Up	2.06E-08	4.05E-08	Uncharacterized protein OS=Anopheles darlingi GN=AND_27065 PE=4 SV=1
CCG021480.1	456	18	82	1.69617373	6.286356788	1.889940207	Up	2.06E-08	4.06E-08	AAEL012377-PA OS=Aedes aegypti GN=AAEL012377 PE=4 SV=1

CCG026343.1	816	20	82	1.05317977	3.512964087	1.737937114	Up	1.10E-07	2.11E-07	AAEL009952-PA OS=Aedes aegypti GN=AAEL009952 PE=4 SV=1
CCG015101.1	1502	22	82	0.6293836	1.908507787	1.60043359	Up	5.43E-07	1.01E-06	AAEL015428-PA OS=Aedes aegypti GN=AAEL015428 PE=4 SV=1
CCG014053.1	252	164	82	27.9644305	11.37531228	-1.297686796	Down	6.13E-12	1.38E-11	AAEL011157-PA OS=Aedes aegypti GN=AAEL011157 PE=4 SV=1
CCG012952.1	1662	198	82	5.11913806	1.724776592	-1.569491412	Down	2.49E-18	9.54E-18	AAEL012412-PA OS=Aedes aegypti GN=AAEL012412 PE=4 SV=1
CCG004692.1	1038	201	82	8.32072896	2.761636508	-1.591186483	Down	6.22E-19	2.40E-18	AAEL007106-PA OS=Aedes aegypti GN=AAEL007106 PE=3 SV=1
CCG006548.1	279	325	82	50.0543504	10.27447561	-2.284430699	Down	2.02E-47	9.32E-47	-
CCG007335.1	1850	647	82	15.0277937	1.549501997	-3.277756694	Down	4.10E-137	2.30E-136	AAEL001818-PA OS=Aedes aegypti GN=AAEL001818 PE=4 SV=1
CCG020771.1	1128	1044	82	39.7698608	2.54129317	-3.968040788	Down	3.88E-259	2.40E-258	AAEL014045-PA OS=Aedes aegypti GN=AAEL014045 PE=4 SV=1
CCG009080.1	1338	5904	82	189.606363	2.142435497	-6.467611798	Down	0	0	Beta-2 tubulin OS=Aedes aegypti GN=B2t PE=3 SV=1
CCG022346.1	1131	6280	82	238.594106	2.534552339	-6.556683635	Down	0	0	AAEL005961-PA OS=Aedes aegypti GN=AAEL005961 PE=3 SV=1
CCG026009.1	294	1	83	0.14615556	9.869173374	6.077352635	Up	0.000587	0.00091842	Putative 8 kDa secreted peptide 8kD-2 OS=Aedes albopictus PE=4 SV=1
CCG028141.1	267	5	83	0.80467668	10.86717967	3.75542454	Up	1.59E-05	2.76E-05	Transporter OS=Aedes aegypti GN=AAEL003624 PE=3 SV=1
CCG019979.1	1830	17	83	0.3991724	1.585539329	1.989889794	Up	8.27E-09	1.65E-08	AAEL005592-PA OS=Aedes aegypti GN=AAEL005592 PE=4 SV=1
CCG025531.1	638	20	83	1.34701362	4.547863593	1.75542454	Up	7.48E-08	1.44E-07	AAEL004617-PA OS=Aedes aegypti GN=AAEL004617 PE=4 SV=1
CCG024165.1	1413	24	83	0.72984687	2.053458579	1.492390134	Up	1.65E-06	3.01E-06	AAEL004614-PA OS=Aedes aegypti GN=AAEL004614 PE=4 SV=1
CCG022871.1	1122	28	83	1.07232849	2.586040082	1.269997713	Up	2.21E-05	3.80E-05	AAEL005950-PA OS=Aedes aegypti GN=CASPS16 PE=3 SV=1
CCG021574.1	1758	140	83	3.42193563	1.650476093	-1.051930382	Down	7.98E-08	1.54E-07	AAEL012956-PA OS=Aedes aegypti GN=AAEL012956 PE=3 SV=1

CCG006469.1	1086	143	83	5.65807739	2.671765168	-1.082518702	Down	2.80E-08	5.49E-08	AAEL011924-PA OS=Aedes aegypti GN=AAEL011924 PE=3 SV=1
CCG007116.1	1605	145	83	3.88200095	1.807811198	-1.102556455	Down	1.37E-08	2.73E-08	AAEL005207-PA OS=Aedes aegypti GN=AAEL005207 PE=4 SV=1
CCG002589.1	1932	149	83	3.31391846	1.501830731	-1.141815885	Down	3.21E-09	6.54E-09	AAEL005247-PA OS=Aedes aegypti GN=AAEL003247 PE=4 SV=2
CCG008587.1	1311	154	83	5.04755082	2.213224235	-1.189433906	Down	4.98E-10	1.04E-09	GJ16356 OS=Drosophila virilis GN=GJ16356 PE=3 SV=1
CCG013491.1	2206	165	83	3.21396474	1.315293278	-1.288969579	Down	6.82E-12	1.53E-11	AAEL014654-PA OS=Aedes aegypti GN=AAEL014654 PE=4 SV=1
CCG006416.1	1593	236	83	6.36588661	1.821429361	-1.805290414	Down	5.40E-26	2.21E-25	AAEL014035-PA OS=Aedes aegypti GN=AAEL014035 PE=4 SV=1
CCG006221.1	1431	238	83	7.14660855	2.027628911	-1.817465128	Down	1.97E-26	8.09E-26	AAEL011784-PA OS=Aedes aegypti GN=AAEL011784 PE=3 SV=1
CCG004343.1	687	246	83	15.3865425	4.223489042	-1.86516187	Down	3.33E-28	1.39E-27	AGAP007127-PA OS=Anopheles gambiae GN=AGAP007127 PE=4 SV=3
CCG012815.1	315	249	83	33.9665521	9.211228483	-1.882649297	Down	7.09E-29	2.97E-28	AGAP007867-PA (Fragment) OS=Anopheles gambiae GN=AGAP007867 PE=4 SV=3
CCG025346.1	1611	301	83	8.02848549	1.801078195	-2.156267042	Down	4.60E-41	2.06E-40	AGAP001084-PA OS=Anopheles gambiae GN=AgaP_AGAP001084 PE=4 SV=5
CCG001105.1	4603	313	83	2.92190461	0.630357804	-2.212666212	Down	5.28E-44	2.40E-43	AAEL007374-PA OS=Aedes aegypti GN=AAEL007374 PE=4 SV=1
CCG008080.1	1690	358	83	9.10246449	1.716885782	-2.406463142	Down	2.24E-55	1.06E-54	Pol polyprotein (Fragment) OS=Anopheles gambiae PE=4 SV=1
CCG017872.1	1722	488	83	12.1772535	1.68498082	-2.853384702	Down	1.64E-90	8.53E-90	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001797 PE=4 SV=1
CCG025898.1	765	493	83	27.6916067	3.792858787	-2.868091201	Down	6.49E-92	3.38E-91	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005987 PE=4 SV=1
CCG007454.1	318	0	84	0.001	9.234261789	13.17278092	Up	4.44E-16	1.65E-15	-

CCG015326.1	498	0	84	0.001	5.896576805	12.52566194	Up	4.44E-16	1.63E-15	AAEL000870-PA OS=Aedes aegypti GN=AAEL000870 PE=4 SV=1
CCG016848.1	1173	0	84	0.001	2.503406009	11.28967657	Up	4.44E-16	1.63E-15	AAEL000858-PA OS=Aedes aegypti GN=AAEL008858 PE=4 SV=1
CCG011666.1	822	3	84	0.15682385	3.572378648	4.509668126	Up	5.28E-05	8.84E-05	AAEL011537-PA OS=Aedes aegypti GN=AAEL011537 PE=4 SV=1
CCG005173.1	528	5	84	0.40691037	5.561544032	3.772702532	Up	1.59E-05	2.76E-05	-
CCG021866.1	1059	8	84	0.32460612	2.772894475	3.094630627	Up	2.19E-06	3.96E-06	AAEL005438-PA OS=Aedes aegypti GN=AAEL005438 PE=4 SV=1
CCG018074.1	383	9	84	1.00973267	7.667089423	2.924705625	Up	1.09E-06	2.01E-06	-
CCG019547.1	336	11	84	1.40674726	8.739569193	2.635199008	Up	2.67E-07	5.03E-07	Uncharacterized protein OS=Anopheles darlingi GN=AND_13030 PE=4 SV=1
CCG003727.1	911	12	84	0.56601187	3.223375685	2.509668126	Up	1.30E-07	2.49E-07	AAEL000891-PA OS=Aedes aegypti GN=AAEL008911 PE=4 SV=1
CCG015701.1	1273	19	84	0.64133932	2.306751963	1.846703113	Up	2.18E-08	4.30E-08	AAEL010089-PA OS=Aedes aegypti GN=AAEL010089 PE=4 SV=1
CCG010287.1	1625	28	84	0.74040158	1.807073999	1.287275705	Up	1.61E-05	2.78E-05	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG024777.1	1440	28	84	0.83552262	2.039232812	1.287275705	Up	1.61E-05	2.78E-05	AAEL002292-PA OS=Aedes aegypti GN=AAEL002292 PE=4 SV=1
CCG000637.1	471	30	84	2.73692577	6.234597131	1.187740031	Up	5.10E-05	8.57E-05	AAEL005713-PA OS=Aedes aegypti GN=AAEL005713 PE=4 SV=1
CCG004733.1	1158	137	84	5.08363872	2.535833548	-1.003401456	Down	3.28E-07	6.16E-07	AAEL008226-PA OS=Aedes aegypti GN=AAEL008226 PE=4 SV=1
CCG020963.1	421	154	84	15.7181452	6.975048097	-1.172155914	Down	7.88E-10	1.63E-09	AAEL008169-PA OS=Aedes aegypti GN=AAEL008169 PE=3 SV=1
CCG009829.2	2892	163	84	2.42187647	1.015385632	-1.254097528	Down	2.49E-11	5.43E-11	AAEL007967-PA OS=Aedes aegypti GN=AAEL007967 PE=4 SV=1
CCG004945.1	3561	175	84	2.11168311	0.824626579	-1.356580485	Down	1.93E-13	5.57E-13	AAEL001584-PA OS=Aedes aegypti GN=AAEL001584 PE=4 SV=1
CCG002559.1	1050	179	84	7.32531666	2.796662142	-1.389185151	Down	3.62E-14	1.25E-13	AAEL007104-PA OS=Aedes aegypti GN=AAEL007104 PE=4 SV=1

CCG004398.1	3980	180	84	1.94335483	0.737812877	-1.39722247	Down	2.37E-14	8.26E-14	Reverse transcriptase homolog protein OS=Chironomus thummi GN=reverse transcriptase homolog PE=4 SV=1
CCG014878.1	2067	246	84	5.11395971	1.42065566	-1.847883879	Down	7.27E-28	3.02E-27	AAEL011763-PA OS=Aedes aegypti GN=PPO3 PE=4 SV=1
CCG016069.1	1371	248	84	7.77278934	2.141863785	-1.859565684	Down	2.61E-28	1.09E-27	AAEL011334-PA OS=Aedes aegypti GN=AAEL011334 PE=4 SV=1
CCG016740.1	1719	276	84	6.8991546	1.708257853	-2.01389383	Down	1.05E-34	4.53E-34	AAEL000933-PA OS=Aedes aegypti GN=AAEL000933 PE=4 SV=1
CCG022033.1	4416	285	84	2.7731826	0.664967221	-2.060187482	Down	7.97E-37	3.49E-36	Leucine-rich repeat-containing G-protein coupled receptor 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ013183 PE=4 SV=1
CCG020587.1	5370	355	84	2.84064354	0.54683338	-2.377044588	Down	3.85E-54	1.82E-53	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015383 PE=4 SV=1
CCG002675.1	1169	380	84	13.9679206	2.511971984	-2.475224982	Down	1.27E-60	6.12E-60	AAEL011200-PA OS=Aedes aegypti GN=AAEL011200 PE=4 SV=1
CCG020913.1	1851	1059	84	24.5839811	1.586437196	-3.953856247	Down	4.09E-262	2.53E-261	AAEL004014-PA OS=Aedes aegypti GN=AAEL004014 PE=3 SV=1
CCG001167.1	11724	1179	84	4.32116318	0.250468718	-4.108717376	Down	4.00E-300	2.52E-299	AAEL014313-PA OS=Aedes aegypti GN=AAEL014313 PE=4 SV=1
CCG010369.1	4110	2657	84	27.7787311	0.71447573	-5.280951886	Down	0	0	AAEL012794-PA OS=Aedes aegypti GN=AAEL012794 PE=4 SV=1
CCG015454.1	852	0	85	0.001	3.487621509	11.76802777	Up	4.44E-16	1.63E-15	AAEL003126-PA OS=Aedes aegypti GN=AAEL003128 PE=4 SV=1
CCG023233.1	885	4	85	0.19421349	3.357574605	4.11170414	Up	2.95E-05	5.03E-05	Serine protease OS=Culex quinquefasciatus GN=CpipJ_CPIJ002753 PE=3 SV=1
CCG026120.1	606	4	85	0.28362861	4.903388656	4.11170414	Up	2.95E-05	5.02E-05	AAEL000163-PA OS=Aedes aegypti GN=AAEL000163 PE=4 SV=1
CCG025807.1	884	6	85	0.29164978	3.361372767	3.526741639	Up	8.38E-06	1.47E-05	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1

CCG002199.1	1257	18	85	0.6153184	2.363924841	1.941779139	Up	7.21E-09	1.44E-08	AAEL008337-PA OS=Aedes aegypti GN=AAEL008337 PE=4 SV=1
CCG017670.1	1127	22	85	0.83880582	2.636604726	1.652272521	Up	1.79E-07	3.39E-07	AAEL011007-PA OS=Aedes aegypti GN=AAEL011007 PE=4 SV=1
CCG007902.1	3464	23	85	0.28530713	0.857809909	1.588142184	Up	3.88E-07	7.26E-07	Pol-like protein OS=Biomphalaria glabrata PE=4 SV=1
CCG000197.1	987	24	85	1.04485677	3.010591211	1.526741639	Up	8.13E-07	1.50E-06	AAEL012538-PA OS=Aedes aegypti GN=AAEL012538 PE=4 SV=1
CCG018723.1	883	25	85	1.21658365	3.365179531	1.46784795	Up	1.65E-06	3.01E-06	-
CCG010910.1	744	33	85	1.90591565	3.993889147	1.067310021	Up	0.0001831	0.00029486	AAEL008376-PA OS=Aedes aegypti GN=AAEL008376 PE=4 SV=1
CCG014844.1	1413	144	85	4.37908123	2.102939509	-1.058220861	Down	4.49E-08	8.74E-08	Sex peptide receptor OS=Aedes aegypti GN=SPR PE=2 SV=1
CCG005884.1	3975	176	85	1.90255932	0.747535478	-1.347727479	Down	2.18E-13	6.26E-13	AAEL012104-PA (Fragment) OS=Aedes aegypti GN=AAEL012104 PE=4 SV=1
CCG009485.1	934	179	85	8.23509903	3.181427758	-1.372111637	Down	6.26E-14	2.07E-13	AAEL012383-PA OS=Aedes aegypti GN=AAEL012383 PE=4 SV=1
CCG019209.1	732	179	85	10.5076264	4.05936274	-1.372111637	Down	6.26E-14	2.07E-13	AAEL002520-PA OS=Aedes aegypti GN=AAEL002520 PE=4 SV=1
CCG003593.1	1803	191	85	4.55197965	1.648060746	-1.465724688	Down	3.64E-16	1.36E-15	AAEL009520-PA OS=Aedes aegypti GN=AAEL009520 PE=4 SV=1
CCG018416.1	2166	195	85	3.86846641	1.3718622	-1.495626174	Down	6.21E-17	2.35E-16	AAEL000474-PA OS=Aedes aegypti GN=AAEL000474 PE=4 SV=1
CCG000809.1	1590	287	85	7.75617222	1.868838695	-2.053202787	Down	6.49E-37	2.85E-36	AAEL014910-PA OS=Aedes aegypti GN=AAEL014910 PE=4 SV=1
CCG010126.1	819	291	85	15.2676346	3.628148383	-2.073171203	Down	7.31E-38	3.22E-37	AAEL002678-PA OS=Aedes aegypti GN=AAEL002678 PE=4 SV=1
CCG001729.1	231	409	85	76.0806124	12.86343518	-2.564252893	Down	8.53E-68	4.21E-67	AAEL000458-PA (Fragment) OS=Aedes aegypti GN=AAEL000458 PE=2 SV=1
CCG024453.1	1419	499	85	15.1105691	2.094047587	-2.851191865	Down	1.94E-92	1.02E-91	Lipase OS=Aedes aegypti GN=AAEL004933 PE=3 SV=1
CCG016362.1	1467	555	85	16.2564436	2.025530692	-3.004639821	Down	2.61E-108	1.40E-107	AAEL004690-PA OS=Aedes aegypti GN=AAEL004690 PE=4 SV=1

CCG016273.1	654	1299	85	85.3481426	4.54350692	-4.231481575	Down	0	0	AAEL013506-PA OS=Aedes aegypti GN=CTLGA2_b PE=4 SV=1
CCG001542.1	1644	1620	85	42.3424392	1.807453483	-4.550073958	Down	0	0	AAEL003038-PA OS=Aedes aegypti GN=AAEL003038 PE=4 SV=1
CCG017210.1	1536	1842	85	51.5301114	1.934540056	-4.735353206	Down	0	0	AAEL004046-PA OS=Aedes aegypti GN=AAEL004048 PE=4 SV=1
CCG016526.1	774	3	86	0.16654936	3.884252975	4.543615458	Up	5.28E-05	8.86E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004372 PE=4 SV=1
CCG021960.1	1332	3	86	0.09677868	2.257065918	4.543615458	Up	5.28E-05	8.86E-05	AAEL003984-PA OS=Aedes aegypti GN=AAEL003984 PE=4 SV=1
CCG008861.1	1068	9	86	0.36210451	2.814992324	2.958652957	Up	1.09E-06	2.01E-06	AAEL011858-PA OS=Aedes aegypti GN=AAEL011858 PE=4 SV=1
CCG010541.1	588	10	86	0.7307778	5.112945242	2.806649864	Up	5.43E-07	1.01E-06	tRNA-dihydrouridine synthase OS=Aedes aegypti GN=AAEL005898 PE=3 SV=1
CCG001909.1	477	13	86	1.17108291	6.30275011	2.42813824	Up	6.34E-08	1.22E-07	AAEL007556-PA OS=Aedes aegypti GN=AAEL007556 PE=4 SV=1
CCG001148.1	999	19	86	0.8172422	3.009421224	1.880650445	Up	1.02E-08	2.04E-08	Guanylate cyclase OS=Aedes aegypti GN=AAEL007359 PE=3 SV=1
CCG007102.1	561	19	86	1.45530295	5.359022821	1.880650445	Up	1.02E-08	2.04E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015728 PE=4 SV=1
CCG016981.1	1573	20	86	0.54634119	1.911259887	1.806649864	Up	2.35E-08	4.62E-08	AAEL010790-PA OS=Aedes aegypti GN=AAEL010790 PE=4 SV=1
CCG002771.1	315	21	86	2.86464897	9.544164452	1.736260536	Up	5.44E-08	1.06E-07	AAEL009672-PA OS=Aedes aegypti GN=AAEL009672 PE=4 SV=1
CCG022658.1	472	30	86	2.7311272	6.369516531	1.221687363	Up	2.77E-05	4.74E-05	-
CCG013819.1	1431	33	86	0.99091631	2.100916703	1.084183839	Up	0.000138	0.00022366	AAEL007305-PA OS=Aedes aegypti GN=AAEL007305 PE=4 SV=1
CCG027467.2	439	34	86	3.32795211	6.848318457	1.041115117	Up	0.0002257	0.00036171	-
CCG022095.1	1308	143	86	4.69776151	2.298479971	-1.031293378	Down	9.45E-08	1.82E-07	AAEL006265-PA (Fragment) OS=Aedes aegypti GN=AAEL006265 PE=4 SV=1

CCG011516.1	1341	150	86	4.80645801	2.241917824	-1.100240732	Down	8.21E-09	1.64E-08	AAEL011899-PA OS=Aedes aegypti GN=AAEL011899 PE=4 SV=1
CCG016757.1	876	151	86	7.40688348	3.431976943	-1.109826781	Down	5.73E-09	1.15E-08	-
CCG024633.1	1398	187	86	5.74773989	2.150509158	-1.418316501	Down	3.67E-15	1.33E-14	AAEL014355-PA OS=Aedes aegypti GN=AAEL014355 PE=4 SV=1
CCG004241.1	796	205	86	11.0663261	3.776899249	-1.550902141	Down	1.27E-18	4.89E-18	AAEL000692-PA OS=Aedes aegypti GN=CYP6AG6 PE=3 SV=1
CCG011445.2	1398	237	86	7.28456874	2.150509158	-1.76016529	Down	3.04E-25	1.24E-24	AAEL014355-PA OS=Aedes aegypti GN=AAEL014355 PE=4 SV=1
CCG027498.1	1218	277	86	9.77226313	2.468318393	-1.985164208	Down	3.38E-34	1.46E-33	AGAP000138-PB OS=Anopheles gambiae GN=AgaP_AGAP000138 PE=2 SV=1
CCG002414.1	777	338	86	18.6921111	3.869255859	-2.272301478	Down	6.16E-49	2.86E-48	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005987 PE=4 SV=1
CCG005177.1	849	368	86	18.6252795	3.541121086	-2.394983998	Down	1.47E-56	7.02E-56	AAEL000870-PA OS=Aedes aegypti GN=AAEL008701 PE=4 SV=1
CCG004658.1	1514	456	86	12.9420073	1.985740953	-2.704312056	Down	5.70E-80	2.90E-79	AAEL0003192-PA OS=Aedes aegypti GN=AAEL003192 PE=4 SV=1
CCG005951.1	1797	862	86	20.6120819	1.673017141	-3.622966101	Down	3.90E-199	2.33E-198	AAEL0006064-PA OS=Aedes aegypti GN=AAEL006064 PE=4 SV=1
CCG005322.1	1223	6620	86	232.591695	2.458227148	-6.564037543	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018952 PE=4 SV=1
CCG000328.1	864	0	87	0.001	3.520104258	11.78140244	Up	4.44E-16	1.63E-15	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008885 PE=4 SV=1
CCG014091.1	544	15	87	1.18482724	5.590753822	2.238366104	Up	1.49E-08	2.94E-08	AAEL004644-PA OS=Aedes aegypti GN=AAEL004644 PE=4 SV=1
CCG021691.1	2093	15	87	0.30795319	1.453115184	2.238366104	Up	1.49E-08	2.94E-08	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG013065.1	3059	18	87	0.25284577	0.994236705	1.975331698	Up	4.07E-09	8.23E-09	Polyprotein OS=Mizuhopecten yessoensis PE=4 SV=1
CCG018742.1	724	24	87	1.42441109	4.200787402	1.560294199	Up	3.97E-07	7.44E-07	AAEL001764-PA OS=Aedes aegypti GN=AAEL001764 PE=4 SV=1

CCG000655.1	1992	27	87	0.5824211	1.526792208	1.390369198	Up	3.22E-06	5.77E-06	AAEL013488-PA OS=Aedes aegypti GN=AAEL013488 PE=3 SV=1
CCG006140.1	1444	29	87	0.86296558	2.106211966	1.287275705	Up	1.13E-05	1.97E-05	LINE-1 retrotransposon-like element, putative (Fragment) OS=Aspergillus flavus (strain ATCC 200026 / FGSC A1120 / NRRL 3357 / JCM 12722 / SRRC 167) GN=AFLA_063980 PE=4 SV=1
CCG020722.1	638	29	87	1.95316976	4.767037742	1.287275705	Up	1.13E-05	1.97E-05	-
CCG022158.1	1587	33	87	0.89351055	1.916427271	1.10086258	Up	0.0001038	0.00016918	AAEL013562-PA OS=Aedes aegypti GN=AAEL013562 PE=4 SV=1
CCG020402.1	498	151	87	13.0289758	6.107168834	-1.09314804	Down	8.77E-09	1.75E-08	AAEL007773-PA OS=Aedes aegypti GN=AAEL007773 PE=4 SV=1
CCG011804.1	552	178	87	13.8561825	5.509728404	-1.330476731	Down	2.76E-13	7.70E-13	AAEL006454-PA OS=Aedes aegypti GN=AAEL006454 PE=4 SV=1
CCG000250.1	2037	191	87	4.02907183	1.493063367	-1.432172128	Down	1.16E-15	4.22E-15	Transmembrane protease OS=Culex quinquefasciatus GN=CpipJ_CPIJ007099 PE=3 SV=1
CCG003964.1	1401	210	87	6.44085958	2.170856588	-1.568988818	Down	2.43E-19	9.41E-19	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001097 PE=4 SV=1
CCG016676.1	747	218	87	12.5400296	4.071445889	-1.622927625	Down	5.90E-21	2.32E-20	AAEL000457-PA OS=Aedes aegypti GN=AAEL000457 PE=4 SV=1
CCG017064.1	948	364	87	16.4989276	3.208196286	-2.362537941	Down	4.51E-55	2.14E-54	AAEL004353-PA OS=Aedes aegypti GN=AAEL004353 PE=4 SV=1
CCG009966.1	807	746	87	39.7217125	3.768736158	-3.397775121	Down	3.87E-163	2.23E-162	Putative secreted protein OS=Aedes aegypti PE=2 SV=1
CCG000284.1	759	6909	87	391.143474	4.007075203	-6.609004497	Down	0	0	AAEL011802-PA OS=Aedes aegypti GN=AAEL011802 PE=4 SV=1
CCG019520.1	1239	2	88	0.06936196	2.482912313	5.161744822	Up	9.36E-05	0.00015345	-
CCG008396.1	378	3	88	0.34102964	8.138434804	4.576782322	Up	5.28E-05	8.85E-05	AAEL004541-PA OS=Aedes aegypti GN=AAEL004541 PE=4 SV=1
CCG014682.1	399	6	88	0.64616142	7.71009613	3.576782322	Up	8.38E-06	1.48E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001479 PE=4 SV=1

CCG023561.1	1033	8	88	0.33277626	2.97805262	3.161744822	Up	2.19E-06	3.96E-06	AAEL002757-PA OS=Aedes aegypti GN=AAEL002757 PE=4 SV=1
CCG010711.1	645	10	88	0.66619744	4.769501327	2.839816728	Up	5.43E-07	1.01E-06	Transporter OS=Aedes aegypti GN=AAEL003624 PE=3 SV=1
CCG004156.1	933	11	88	0.50660995	3.297243683	2.702313204	Up	2.67E-07	5.03E-07	AAEL004637-PA OS=Aedes aegypti GN=AAEL004837 PE=4 SV=1
CCG008483.1	1011	17	88	0.72253758	3.04285693	2.074281981	Up	3.99E-09	8.06E-09	AAEL007064-PA OS=Aedes aegypti GN=GNBPB6 PE=2 SV=1
CCG020113.1	1134	24	88	0.90941237	2.712811601	1.576782322	Up	2.77E-07	5.20E-07	AAEL015081-PA OS=Aedes aegypti GN=AAEL015081 PE=3 SV=1
CCG013536.1	1422	27	88	0.81588104	2.163381404	1.40685732	Up	2.30E-06	4.15E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002266 PE=4 SV=1
CCG009064.1	1482	30	88	0.86983268	2.075795112	1.254854227	Up	1.49E-05	2.59E-05	AAEL007260-PA OS=Aedes aegypti GN=AAEL007260 PE=4 SV=1
CCG014773.1	3036	30	88	0.42460212	1.013283385	1.254854227	Up	1.49E-05	2.59E-05	AAEL012498-PA OS=Aedes aegypti GN=AAEL012498 PE=4 SV=1
CCG009849.1	813	31	88	1.63845237	3.783921717	1.207548512	Up	2.64E-05	4.52E-05	AAEL000174-PA OS=Aedes aegypti GN=AAEL000174 PE=4 SV=1
CCG006023.1	920	35	88	1.63471816	3.34383517	1.032461806	Up	0.0002102	0.00033747	-
CCG021398.1	945	185	88	8.41206445	3.255373922	-1.369636638	Down	2.60E-14	9.05E-14	AAEL010068-PA OS=Aedes aegypti GN=AAEL010068 PE=4 SV=1
CCG026713.1	1428	197	88	5.92789756	2.154291566	-1.460306997	Down	1.53E-16	5.74E-16	-
CCG014119.1	10561	215	88	0.87477445	0.291291389	-1.586448027	Down	4.60E-20	1.80E-19	AAEL009419-PA (Fragment) OS=Aedes aegypti GN=AAEL009419 PE=4 SV=1
CCG002398.1	1659	226	88	5.85362268	1.854326917	-1.65843414	Down	2.62E-22	1.04E-21	AAEL000804-PA OS=Aedes aegypti GN=AAEL000804 PE=4 SV=1
CCG019949.1	1587	232	88	6.28164992	1.938455171	-1.696236173	Down	1.47E-23	5.91E-23	AAEL014910-PA OS=Aedes aegypti GN=AAEL014910 PE=4 SV=1
CCG025992.1	1233	282	88	9.82762787	2.494994611	-1.97780653	Down	1.28E-34	5.52E-34	AAEL009864-PA OS=Aedes aegypti GN=AAEL009864 PE=4 SV=1
CCG003787.1	762	289	88	16.2969203	4.03717632	-2.01318086	Down	3.01E-36	1.31E-35	-

CCG021805.1	1854	433	88	10.0355421	1.659292533	-2.596478392	Down	1.10E-72	5.49E-72	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005453 PE=4 SV=1
CCG013056.1	381	736	88	83.0071514	8.07435264	-3.361817134	Down	2.00E-159	1.15E-158	AAEL002920-PA OS=Aedes aegypti GN=AAEL002920 PE=4 SV=1
CCG012756.1	477	991	88	89.2725514	6.449325694	-3.790996425	Down	2.06E-237	1.26E-236	AAEL008764-PA OS=Aedes aegypti GN=AAEL008764 PE=4 SV=1
CCG000991.1	579	5	89	0.37106852	5.373552043	3.85611854	Up	1.59E-05	2.76E-05	AAEL008915-PA OS=Aedes aegypti GN=AAEL008915 PE=4 SV=1
CCG007006.1	504	20	89	1.7051482	6.173187763	1.85611854	Up	7.39E-09	1.48E-08	AAEL011142-PA OS=Aedes aegypti GN=AAEL011142 PE=4 SV=1
CCG018482.1	480	22	89	1.96944617	6.481847152	1.718615016	Up	3.95E-08	7.71E-08	AAEL006177-PA OS=Aedes aegypti GN=AAEL006177 PE=4 SV=1
CCG006575.1	537	28	89	2.24050758	5.793829856	1.370691713	Up	3.16E-06	5.67E-06	AAEL005735-PA OS=Aedes aegypti GN=AAEL005735 PE=4 SV=1
CCG016898.1	1395	152	89	4.68200693	2.230312998	-1.069880879	Down	1.42E-08	2.81E-08	AAEL007606-PA OS=Aedes aegypti GN=CYP4D39 PE=3 SV=1
CCG026866.1	4210	166	89	1.69429357	0.739022953	-1.196992797	Down	8.54E-11	1.83E-10	-
CCG002693.1	1017	243	89	10.2671047	3.059278892	-1.746765869	Down	1.39E-25	5.69E-25	Bruchpilot, isoform G OS=Drosophila melanogaster GN=brp PE=4 SV=2
CCG016843.1	1119	296	89	11.3664356	2.780417009	-2.031406731	Down	1.64E-37	7.21E-37	AAEL009635-PA OS=Aedes aegypti GN=AAEL009635 PE=4 SV=1
CCG016119.1	11883	1182	89	4.27419223	0.261826696	-4.028967685	Down	1.51E-296	9.47E-296	AAEL000512-PA OS=Aedes aegypti GN=AAEL000512 PE=4 SV=1
CCG019884.1	1461	1493	89	43.9108924	2.129559639	-4.365951815	Down	0	0	AAEL008062-PA (Fragment) OS=Aedes aegypti GN=AAEL008062 PE=4 SV=1
CCG014029.1	1110	17210	89	666.224444	2.80296093	-7.892912842	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_01528 PE=4 SV=1
CCG019189.1	312	4	90	0.55089403	10.0841183	4.1941663	Up	2.95E-05	5.03E-05	Synaptic vesicular amine transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ012480 PE=4 SV=1

CCG005276.1	1970	7	90	0.15268434	1.597078634	3.386811378	Up	4.31E-06	7.69E-06	BEL12_AG transposon polyprotein OS=Anopheles gambiae GN=BEL12-AG transposon PE=4 SV=1
CCG027342.1	519	10	90	0.82793323	6.06212892	2.872238205	Up	5.43E-07	1.01E-06	AAEL000267-PA OS=Aedes aegypti GN=AAEL000267 PE=4 SV=1
CCG025976.1	441	12	90	1.16924448	7.134342198	2.609203799	Up	1.30E-07	2.49E-07	AAEL004210-PA OS=Aedes aegypti GN=AAEL004210 PE=4 SV=1
CCG007223.1	1224	13	90	0.4563779	2.570461527	2.493726582	Up	6.34E-08	1.22E-07	-
CCG007720.1	1077	24	90	0.95754283	2.921304466	1.609203799	Up	1.33E-07	2.54E-07	AAEL009129-PA OS=Aedes aegypti GN=CYP6Z9 PE=3 SV=1
CCG022186.1	1236	26	90	0.90389409	2.54550559	1.493726582	Up	5.82E-07	1.08E-06	AAEL001925-PA OS=Aedes aegypti GN=AAEL001925 PE=4 SV=1
CCG004450.1	3522	29	90	0.35381099	0.893312013	1.336185305	Up	4.29E-06	7.67E-06	AAEL007304-PA OS=Aedes aegypti GN=AAEL007304 PE=4 SV=1
CCG022996.1	1392	31	90	0.95694093	2.260233412	1.23996999	Up	1.43E-05	2.48E-05	AAEL005058-PA OS=Aedes aegypti GN=AAEL005058 PE=4 SV=1
CCG004079.1	1359	256	90	8.09437238	2.315117667	-1.8058337	Down	4.49E-28	1.87E-27	AAEL012720-PA OS=Aedes aegypti GN=AAEL012720 PE=4 SV=1
CCG019191.1	585	272	90	19.9790903	5.378196427	-1.893296541	Down	1.26E-31	5.37E-31	Calcium-activated potassium channel alpha chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ005769 PE=4 SV=1
CCG026366.1	566	340	90	25.8122081	5.558736589	-2.215224636	Down	1.01E-47	4.65E-47	-
CCG017978.1	918	346	90	16.1955645	3.427282036	-2.240461927	Down	3.29E-49	1.53E-48	AAEL004096-PA OS=Aedes aegypti GN=AAEL004096 PE=4 SV=1
CCG014455.1	2148	413	90	8.26187169	1.464732267	-2.495831671	Down	2.27E-66	1.11E-65	AAEL009203-PA OS=Aedes aegypti GN=AAEL009203 PE=3 SV=1
CCG008805.1	931	599	90	27.646478	3.379425252	-3.032245893	Down	7.77E-118	4.24E-117	AAEL009295-PA OS=Aedes aegypti GN=AAEL009295 PE=4 SV=1
CCG018448.1	852	674	90	33.9924896	3.692775715	-3.202438481	Down	1.19E-139	6.69E-139	AAEL009296-PA OS=Aedes aegypti GN=AAEL009298 PE=4 SV=1
CCG027571.1	426	4	91	0.40347169	7.467613113	4.210107844	Up	2.95E-05	5.04E-05	AAEL002491-PA OS=Aedes aegypti GN=AAEL002491 PE=4 SV=1

CCG011638.1	1631	10	91	0.26345637	1.950461794	2.888179749	Up	5.43E-07	1.01E-06	AAEL010625-PA OS=Aedes aegypti GN=AAEL010625 PE=4 SV=1
CCG001366.1	483	16	91	1.42342806	6.586342001	2.210107844	Up	7.15E-09	1.43E-08	AAEL004301-PA OS=Aedes aegypti GN=AAEL004301 PE=2 SV=1
CCG004647.1	552	17	91	1.32334328	5.763049251	2.122645003	Up	3.56E-09	7.21E-09	AAEL007970-PA OS=Aedes aegypti GN=AAEL007970 PE=4 SV=1
CCG008402.1	984	32	91	1.39738974	3.232930067	1.210107844	Up	1.86E-05	3.20E-05	AAEL004589-PA OS=Aedes aegypti GN=AAEL004589 PE=3 SV=1
CCG003253.1	2469	34	91	0.59172579	1.288458156	1.122645003	Up	5.48E-05	9.13E-05	AAEL002126-PA OS=Aedes aegypti GN=CLIPA15 PE=3 SV=2
CCG026939.1	1100	156	91	6.09388964	2.892002897	-1.075294375	Down	7.95E-09	1.59E-08	faty acid desaturase OS=Culex quinquefasciatus GN=CpipJ_CPIJ004097 PE=4 SV=1
CCG000861.1	1515	157	91	4.4529692	2.099804083	-1.084512905	Down	5.59E-09	1.13E-08	Cytochrome P450 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005957 PE=3 SV=1
CCG028500.2	993	171	91	7.39962197	3.203628586	-1.207744671	Down	3.24E-11	7.03E-11	AAEL000574-PA OS=Aedes aegypti GN=AAEL000574 PE=4 SV=1
CCG001155.1	966	182	91	8.0957471	3.293171	-1.297686796	Down	4.32E-13	1.14E-12	AAEL008219-PA OS=Aedes aegypti GN=AAEL008219 PE=4 SV=1
CCG010608.2	1156	218	91	8.10328905	2.75190587	-1.558076481	Down	7.88E-20	3.07E-19	Guanylate cyclase OS=Aedes aegypti GN=AAEL006806 PE=3 SV=1
CCG016719.1	1908	258	91	5.81037292	1.667297267	-1.801119411	Down	3.49E-28	1.45E-27	AAEL007378-PA OS=Aedes aegypti GN=AAEL007378 PE=4 SV=1
CCG024778.1	963	366	91	16.3311764	3.3034301	-2.305591994	Down	8.78E-54	4.15E-53	inositol polyphosphate 1-phosphatase OS=Culex quinquefasciatus GN=CpipJ_CPIJ015548 PE=4 SV=1
CCG028323.1	1012	7	92	0.29722148	3.178025161	3.418520238	Up	4.31E-06	7.69E-06	AAEL015563-PA (Fragment) OS=Aedes aegypti GN=AAEL015563 PE=3 SV=1
CCG014920.1	993	15	92	0.64908965	3.238833296	2.318984564	Up	1.48E-08	2.92E-08	AAEL000409-PA OS=Aedes aegypti GN=AAEL000409 PE=4 SV=1

CCG004479.1	1092	18	92	0.70829233	2.945202805	2.055950158	Up	1.92E-09	3.94E-09	AAEL002106-PA OS=Aedes aegypti GN=AAEL002106 PE=4 SV=1
CCG006910.1	1251	24	92	0.82435942	2.570872472	1.640912659	Up	6.37E-08	1.23E-07	AAEL001631-PA OS=Aedes aegypti GN=AAEL001631 PE=4 SV=1
CCG000444.1	624	30	92	2.06585263	5.154104909	1.318984564	Up	4.17E-06	7.46E-06	Peptidyl-prolyl cis-trans isomerase OS=Aedes aegypti GN=AAEL011758 PE=3 SV=1
CCG013398.1	720	33	92	1.96944617	4.466890921	1.181481041	Up	2.39E-05	4.10E-05	AAEL011715-PA OS=Aedes aegypti GN=AAEL011715 PE=4 SV=1
CCG015035.1	1197	33	92	1.18462927	2.686851682	1.181481041	Up	2.39E-05	4.10E-05	AAEL003024-PA OS=Aedes aegypti GN=AAEL003024 PE=4 SV=1
CCG005550.1	1593	37	92	0.99804154	2.01893375	1.016421794	Up	0.0001819	0.00029306	AAEL001807-PA OS=Aedes aegypti GN=CYP9M9 PE=3 SV=2
CCG013940.1	255	197	92	33.1962263	12.61239789	-1.39617666	Down	1.50E-15	5.43E-15	AAEL006510-PA (Fragment) OS=Aedes aegypti GN=AAEL006510 PE=4 SV=1
CCG017425.1	1137	207	92	7.82298598	2.82863805	-1.467611798	Down	2.00E-17	7.61E-17	Peroxidasin OS=Aedes aegypti PE=2 SV=1
CCG019907.1	591	344	92	25.0111484	5.441897569	-2.200389595	Down	7.18E-48	3.32E-47	GD19676 OS=Drosophila simulans GN=GD19676 PE=4 SV=1
CCG006995.1	474	4	93	0.36261379	6.858902405	4.241472015	Up	2.95E-05	5.04E-05	AAEL006900-PA OS=Aedes aegypti GN=AAEL006900 PE=4 SV=1
CCG011497.1	859	10	93	0.50022974	3.784772689	2.91954392	Up	5.43E-07	1.01E-06	Putative uncharacterized protein OS=Daphnia pulex GN=DAPPUDRAFT_306575 PE=4 SV=1
CCG008647.1	882	19	93	0.92565188	3.686076803	1.993544501	Up	1.31E-09	2.70E-09	AAEL004482-PA OS=Aedes aegypti GN=AAEL004482 PE=4 SV=1
CCG004646.1	789	24	93	1.30706417	4.120557338	1.656509514	Up	4.39E-08	8.56E-08	AAEL007970-PA OS=Aedes aegypti GN=AAEL007970 PE=4 SV=1
CCG007342.1	820	30	93	1.57206346	3.964780171	1.334581419	Up	3.01E-06	5.42E-06	AAEL009058-PA OS=Aedes aegypti GN=AAEL009058 PE=4 SV=1
CCG026787.1	1697	31	93	0.78495096	1.915804207	1.287275705	Up	5.56E-06	9.87E-06	AAEL014133-PA OS=Aedes aegypti GN=AAEL014133 PE=4 SV=1

CCG013915.1	357	172	93	20.7025052	9.106777983	-1.18479274	Down	5.62E-11	1.21E-10	AAEL002817-PA OS=Aedes aegypti GN=AAEL002817 PE=4 SV=1
CCG001072.1	2688	427	93	6.82592138	1.209493951	-2.496620245	Down	1.38E-68	6.83E-68	AAEL005429-PA OS=Aedes aegypti GN=AAEL005429 PE=4 SV=1
CCG019525.1	679	460	93	29.1105713	4.788099764	-2.604018036	Down	2.33E-77	1.17E-76	-
CCG012509.1	2187	493	93	9.6863645	1.486565953	-2.703971821	Down	2.64E-86	1.36E-85	AAEL010917-PA OS=Aedes aegypti GN=AAEL010917 PE=4 SV=1
CCG015868.1	753	978	93	55.8092967	4.317556095	-3.69221864	Down	2.63E-229	1.60E-228	AAEL006131-PA OS=Aedes aegypti GN=AAEL006131 PE=4 SV=1
CCG018642.1	1386	1207	93	37.4202523	2.345685238	-3.995737946	Down	6.76E-301	4.26E-300	AAEL005917-PA OS=Aedes aegypti GN=AAEL005917 PE=4 SV=1
CCG001143.1	663	10543	93	683.303035	4.903649683	-7.122525807	Down	0	0	AAEL003041-PA OS=Aedes aegypti GN=AAEL003041 PE=4 SV=1
CCG019969.1	360	0	94	0.001	9.127994491	13.15608221	Up	4.44E-16	1.65E-15	AAEL010787-PA OS=Aedes aegypti GN=AAEL010787 PE=3 SV=1
CCG021098.1	491	0	94	0.001	6.692623252	12.70835609	Up	4.44E-16	1.64E-15	-
CCG011738.1	846	0	94	0.001	3.884252975	11.92342145	Up	4.44E-16	1.66E-15	AAEL000769-PA OS=Aedes aegypti GN=AAEL000769 PE=4 SV=1
CCG007699.1	1743	0	94	0.001	1.885300067	10.88057845	Up	4.44E-16	1.64E-15	AAEL013916-PA OS=Aedes aegypti GN=AAEL013916 PE=4 SV=1
CCG014248.1	2184	7	94	0.13772351	1.504614476	3.449547133	Up	4.31E-06	7.69E-06	AAEL014398-PA OS=Aedes aegypti GN=AAEL014398 PE=4 SV=1
CCG001889.1	1234	8	94	0.27857202	2.66294815	3.256902055	Up	2.19E-06	3.96E-06	AAEL006225-PA OS=Aedes aegypti GN=AAEL006225 PE=4 SV=1
CCG024761.2	1302	19	94	0.6270545	2.523869444	2.008974542	Up	1.13E-09	2.33E-09	AAEL006450-PA OS=Aedes aegypti GN=AAEL006450 PE=4 SV=1
CCG021849.2	1650	23	94	0.59897206	1.991562434	1.733340099	Up	1.34E-08	2.66E-08	AAEL004794-PA (Fragment) OS=Aedes aegypti GN=AAEL004794 PE=4 SV=1
CCG027307.1	2975	25	94	0.36109021	1.104564039	1.613045866	Up	6.62E-08	1.28E-07	Uncharacterized protein OS=Anopheles darlingi GN=AND_03682 PE=4 SV=1
CCG009909.1	1341	37	94	1.18559298	2.45046832	1.04744869	Up	0.0001053	0.00017161	AAEL010287-PA OS=Aedes aegypti GN=AAEL010287 PE=4 SV=1

CCG026248.1	741	38	94	2.20357613	4.434653194	1.008974542	Up	0.0001691	0.00027257	AAEL006882-PA OS=Aedes aegypti GN=AAEL006882 PE=4 SV=1
CCG022826.1	454	180	94	17.0364587	7.238057305	-1.234951041	Down	4.22E-12	9.74E-12	-
CCG027378.1	1008	191	94	8.14208265	3.259998032	-1.320526773	Down	5.31E-14	1.78E-13	-
CCG003496.1	1626	204	94	5.39103681	2.02095819	-1.415523286	Down	2.33E-16	8.73E-16	AAEL003960-PA OS=Aedes aegypti GN=AAEL003960 PE=4 SV=1
CCG003291.1	1388	219	94	6.77980683	2.367491366	-1.517885004	Down	3.24E-19	1.25E-18	AAEL004097-PA OS=Aedes aegypti GN=AAEL004097 PE=4 SV=1
CCG022168.1	2724	322	94	5.0793886	1.206342884	-2.074014823	Down	9.66E-42	4.34E-41	AAEL002511-PA (Fragment) OS=Aedes aegypti GN=AAEL002511 PE=4 SV=1
CCG018658.1	898	832	94	39.8116027	3.65932964	-3.443537663	Down	6.11E-184	3.59E-183	-
CCG018907.1	783	1064	94	58.3905461	4.196779076	-3.79838038	Down	3.65E-255	2.25E-254	AAEL007416-PA OS=Aedes aegypti GN=AAEL007416 PE=4 SV=1
CCG027550.2	888	2	95	0.09677868	3.739905736	5.272168812	Up	9.36E-05	0.00015337	AAEL008084-PA OS=Aedes aegypti GN=AAEL008084 PE=4 SV=1
CCG020853.1	972	2	95	0.08841509	3.416704006	5.272168812	Up	9.36E-05	0.00015352	AAEL009134-PA (Fragment) OS=Aedes aegypti GN=AAEL009134 PE=4 SV=1
CCG011424.1	706	5	95	0.30431823	4.704017413	3.950240717	Up	1.59E-05	2.76E-05	-
CCG004529.1	885	10	95	0.48553372	3.752583382	2.950240717	Up	5.43E-07	1.01E-06	AAEL003102-PA OS=Aedes aegypti GN=AAEL003102 PE=3 SV=1
CCG020391.1	471	10	95	0.91230859	7.05103247	2.950240717	Up	5.43E-07	1.01E-06	AAEL007313-PA OS=Aedes aegypti GN=AAEL007313 PE=4 SV=1
CCG020720.1	2178	10	95	0.19728987	1.524810052	2.950240717	Up	5.43E-07	1.01E-06	-
CCG007926.1	1056	12	95	0.48829244	3.144920732	2.687206311	Up	1.30E-07	2.49E-07	AAEL010930-PA OS=Aedes aegypti GN=AAEL010930 PE=4 SV=1
CCG012730.1	849	18	95	0.91101911	3.911703526	2.102243811	Up	1.71E-09	3.49E-09	AAEL008860-PA OS=Aedes aegypti GN=AAEL008860 PE=4 SV=1
CCG025686.1	1161	21	95	0.77723034	2.860496377	1.879851389	Up	1.75E-09	3.58E-09	AAEL011413-PA OS=Aedes aegypti GN=AAEL011413 PE=4 SV=1
CCG001327.1	753	192	95	10.9564264	4.410406764	-1.312793689	Down	5.94E-14	1.98E-13	AAEL010671-PA OS=Aedes aegypti GN=AAEL010671 PE=3 SV=1
CCG011092.1	2214	195	95	3.78459722	1.500016393	-1.335161502	Down	1.75E-14	6.16E-14	AAEL012091-PA OS=Aedes aegypti GN=AAEL012091 PE=3 SV=1

CCG017598.1	411	214	95	22.3735358	8.080380276	-1.469298174	Down	5.44E-18	2.08E-17	histone H2B OS=Aspergillus flavus (strain ATCC 200026 / FGSC A1120 / NRRL 3357 / JCM 12722 / SRR167)
CCG027557.1	2550	324	95	5.45968393	1.302367174	-2.067681191	Down	7.89E-42	3.55E-41	AAEL015663-PA (Fragment) OS=Aedes aegypti GN=AAEL015663 PE=3 SV=1
CCG006077.1	728	403	95	23.7868174	4.56186304	-2.382467216	Down	2.14E-61	1.04E-60	AAEL007054-PA OS=Aedes aegypti GN=AAEL007054 PE=4 SV=1
CCG023063.1	1743	1335	95	32.9114146	1.905356451	-4.110455214	Down	0	0	AAEL004341-PA (Fragment) OS=Aedes aegypti GN=AAEL004341 PE=4 SV=1
CCG006431.1	378	1481	95	168.354966	8.7858103	-4.260187113	Down	0	0	AAEL009337-PA OS=Aedes aegypti GN=AAEL009337 PE=3 SV=1
CCG006787.1	1290	1815	95	60.4574173	2.574446739	-4.553585021	Down	0	0	C(2) long form OS=Culex quinquefasciatus GN=CpipJ_CPIJ017150 PE=4 SV=1
CCG004420.1	477	0	96	0.001	7.03562803	12.78046349	Up	4.44E-16	1.64E-15	AAEL003584-PA OS=Aedes aegypti GN=AAEL003584 PE=4 SV=1
CCG000295.1	768	1	96	0.05595018	4.369784597	6.287275705	Up	0.000587	0.00091717	AAEL004467-PA OS=Aedes aegypti GN=AAEL004467 PE=2 SV=1
CCG020083.1	690	12	96	0.74729973	4.863760247	2.702313204	Up	1.30E-07	2.49E-07	AAEL011163-PA OS=Aedes aegypti GN=AAEL011163 PE=4 SV=1
CCG018444.1	1752	12	96	0.29431325	1.915522015	2.702313204	Up	1.30E-07	2.49E-07	AAEL004778-PA OS=Aedes aegypti GN=AAEL004778 PE=3 SV=1
CCG025939.1	951	14	96	0.63257233	3.52891122	2.479920782	Up	3.07E-08	6.00E-08	AAEL011301-PA OS=Aedes aegypti GN=AAEL011301 PE=4 SV=1
CCG004249.1	1524	19	96	0.53571191	2.202096175	2.039348191	Up	9.27E-10	1.91E-09	AAEL011571-PA (Fragment) OS=Aedes aegypti GN=GPRgr14 PE=3 SV=2
CCG014086.1	1977	21	96	0.45643117	1.697518751	1.894958282	Up	1.23E-09	2.53E-09	zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018423 PE=4 SV=1
CCG010507.1	507	22	96	1.86456442	6.619318679	1.827844086	Up	2.68E-09	5.46E-09	AAEL010139-PA OS=Aedes aegypti GN=AAEL010139 PE=4 SV=1

CCG017837.1	2232	23	96	0.44278848	1.503581797	1.763713748	Up	6.21E-09	1.25E-08	AGAP003323-PA OS=Anopheles gambiae GN=AgaP_AGAP003323 PE=3 SV=5
CCG026521.1	1872	27	96	0.61975579	1.792732142	1.532388202	Up	1.43E-07	2.73E-07	AAEL006738-PA (Fragment) OS=Aedes aegypti GN=AAEL006738 PE=4 SV=1
CCG004164.1	453	29	96	2.75082186	7.408376535	1.429294709	Up	5.80E-07	1.08E-06	AAEL013800-PA OS=Aedes aegypti GN=AAEL013800 PE=4 SV=1
CCG027974.2	1050	31	96	1.26863026	3.196185305	1.333079394	Up	2.12E-06	3.85E-06	AAEL013490-PA OS=Aedes aegypti GN=AAEL013490 PE=4 SV=1
CCG021926.1	222	35	96	6.77450771	15.11709266	1.157992688	Up	2.15E-05	3.69E-05	AAEL000284-PA OS=Aedes aegypti GN=AAEL000284 PE=4 SV=1
CCG019392.1	663	37	96	2.39800932	5.061831931	1.077822339	Up	6.03E-05	0.00010025	AAEL007357-PA OS=Aedes aegypti GN=AAEL007357 PE=4 SV=1
CCG025199.1	345	37	96	4.60834835	9.727520493	1.077822339	Up	6.03E-05	0.00010026	-
CCG021354.1	6750	218	96	1.38776328	0.497184381	-1.48090862	Down	1.69E-18	6.49E-18	Tyrosine-protein kinase receptor (Fragment) OS=Aedes aegypti GN=AAEL007742 PE=3 SV=1
CCG003817.1	381	264	96	29.7743043	8.808384699	-1.757118415	Down	7.12E-28	2.96E-27	AAEL013517-PA OS=Aedes aegypti GN=AAEL013517 PE=4 SV=1
CCG022483.1	1356	284	96	8.99956094	2.474922249	-1.862471415	Down	2.98E-32	1.27E-31	CG34109 OS=Drosophila melanogaster GN=CG12441 PE=2 SV=1
CCG006879.1	1336	394	96	12.6722122	2.511971984	-2.334776115	Down	1.21E-58	5.82E-58	AAEL007363-PA OS=Aedes aegypti GN=AAEL007363 PE=4 SV=1
CCG024180.1	1698	448	96	11.3371267	1.976439676	-2.520079218	Down	1.20E-72	5.99E-72	AAEL003295-PA OS=Aedes aegypti GN=AAEL003295 PE=4 SV=1
CCG004774.1	12210	927	96	3.26232137	0.27485623	-3.569149824	Down	2.74E-211	1.65E-210	AAEL010965-PA OS=Aedes aegypti GN=AAEL010965 PE=4 SV=1
CCG012510.1	2274	944	96	17.8379197	1.475811157	-3.595367345	Down	1.73E-216	1.05E-215	AAEL010921-PA OS=Aedes aegypti GN=AAEL010921 PE=4 SV=1
CCG021497.1	198	3	97	0.65105659	17.12602448	4.717263545	Up	5.28E-05	8.86E-05	-
CCG006525.1	699	4	97	0.24589262	4.851148565	4.302226046	Up	2.95E-05	5.03E-05	AAEL009218-PA OS=Aedes aegypti GN=AAEL009218 PE=3 SV=1

CCG017977.1	828	6	97	0.31137489	4.09535368	3.717263545	Up	8.38E-06	1.48E-05	AAEL004104-PA OS=Aedes aegypti GN=AAEL004104 PE=4 SV=1
CCG005945.1	911	16	97	0.7546825	3.722231446	2.302226046	Up	7.09E-09	1.42E-08	AAEL002864-PA OS=Aedes aegypti GN=AAEL002864 PE=4 SV=1
CCG027260.1	789	25	97	1.36152518	4.297785611	1.658369856	Up	2.18E-08	4.29E-08	AAEL001300-PA OS=Aedes aegypti GN=AAEL001300 PE=4 SV=1
CCG008404.1	1464	27	97	0.79247461	2.316224622	1.547338544	Up	1.00E-07	1.92E-07	AAEL004592-PA OS=Aedes aegypti GN=AAEL004592 PE=4 SV=1
CCG007741.1	3610	27	97	0.32138029	0.939322118	1.547338544	Up	1.00E-07	1.92E-07	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC009401 PE=4 SV=1
CCG022108.2	960	29	97	1.29804407	3.532242549	1.444245051	Up	4.12E-07	7.71E-07	AAEL012479-PA OS=Aedes aegypti GN=AAEL012479 PE=4 SV=1
CCG023248.1	729	29	97	1.70935844	4.651512822	1.444245051	Up	4.12E-07	7.71E-07	Lupus rabnonucleoprotein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008950 PE=4 SV=1
CCG005198.1	828	30	97	1.55687444	4.09535368	1.39533545	Up	8.04E-07	1.49E-06	AAEL006410-PA OS=Aedes aegypti GN=AAEL006410 PE=4 SV=1
CCG026298.1	618	36	97	2.50309134	5.486978717	1.132301045	Up	2.72E-05	4.65E-05	AAEL013310-PA OS=Aedes aegypti GN=AAEL013310 PE=4 SV=1
CCG025114.1	861	38	97	1.89645751	3.938388905	1.054298533	Up	7.45E-05	0.00012341	AAEL003451-PA OS=Aedes aegypti GN=AAEL003451 PE=4 SV=1
CCG014777.2	597	176	97	12.6677945	5.679988018	-1.157205573	Down	7.76E-11	1.67E-10	AAEL012401-PA OS=Aedes aegypti GN=AAEL012401 PE=3 SV=1
CCG020894.1	637	196	97	13.2214568	5.323316871	-1.312483798	Down	3.30E-14	1.15E-13	AAEL000644-PA OS=Aedes aegypti GN=AAEL000644 PE=4 SV=1
CCG019222.1	933	218	97	10.040088	3.634461787	-1.465958279	Down	3.05E-18	1.17E-17	AAEL001338-PA OS=Aedes aegypti GN=AAEL001338 PE=4 SV=1
CCG004072.1	1131	499	97	18.9583533	2.998189962	-2.660669959	Down	8.44E-86	4.34E-85	AAEL013372-PA OS=Aedes aegypti GN=AAEL013372 PE=3 SV=1
CCG013757.1	1398	506	97	15.5527079	2.425574282	-2.680767529	Down	1.06E-87	5.46E-87	AAEL005261-PA OS=Aedes aegypti GN=AAEL005261 PE=4 SV=1

CCG018126.1	528	0	98	0.001	6.488468037	12.66366217	Up	4.44E-16	1.63E-15	60S ribosomal protein L17 OS=Culex quinquefasciatus GN=CpipJ_CPIJ015624 PE=3 SV=1
CCG025877.1	486	2	98	0.17683018	7.049199843	5.317023048	Up	9.36E-05	0.00015338	AAEL009296-PA OS=Aedes aegypti GN=AAEL009296 PE=4 SV=1
CCG025157.1	633	2	98	0.13576535	5.41218187	5.317023048	Up	9.36E-05	0.00015386	AGAP010064-PA OS=Anopheles gambiae GN=AGAP010064 PE=4 SV=1
CCG026742.1	984	15	98	0.65502644	3.481616996	2.410132452	Up	1.48E-08	2.92E-08	AAEL008970-PA OS=Aedes aegypti GN=AAEL008970 PE=4 SV=1
CCG027744.1	378	15	98	1.7051482	9.063256941	2.410132452	Up	1.48E-08	2.92E-08	AAEL009836-PA OS=Aedes aegypti GN=AAEL009836 PE=4 SV=1
CCG003357.1	2136	16	98	0.32187067	1.603890976	2.317023048	Up	7.09E-09	1.42E-08	N-acetylglucosaminyltransferase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005697 PE=3 SV=1
CCG028571.1	1821	16	98	0.37754846	1.881335049	2.317023048	Up	7.09E-09	1.42E-08	AAEL006453-PA OS=Aedes aegypti GN=AAEL006453 PE=4 SV=1
CCG001908.1	420	26	98	2.66003119	8.156931247	1.61658333	Up	3.29E-08	6.43E-08	AAEL007556-PA OS=Aedes aegypti GN=AAEL007556 PE=4 SV=1
CCG028531.1	2480	28	98	0.48514216	1.381415776	1.509668126	Up	1.45E-07	2.76E-07	GK25464 OS=Drosophila willistoni GN=GK25464 PE=4 SV=1
CCG017581.1	216	36	98	7.16162244	15.86069965	1.147098046	Up	2.03E-05	3.49E-05	AAEL002121-PA OS=Aedes aegypti GN=AAEL002121 PE=4 SV=1
CCG011130.1	1393	37	98	1.14133538	2.459376255	1.107569682	Up	3.42E-05	5.78E-05	AAEL003776-PA OS=Aedes aegypti GN=AAEL003776 PE=4 SV=1
CCG028433.1	1722	164	98	4.09235568	1.989495426	-1.040528957	Down	8.50E-09	1.70E-08	AAEL005440-PA OS=Aedes aegypti GN=AAEL005440 PE=4 SV=1
CCG003966.1	1689	214	98	5.44435951	2.028366562	-1.424443938	Down	3.09E-17	1.17E-16	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003199 PE=4 SV=1
CCG011736.1	1689	293	98	7.45419316	2.028366562	-1.877733806	Down	1.41E-33	6.04E-33	AAEL006905-PA OS=Aedes aegypti GN=AAEL006905 PE=4 SV=1
CCG019211.1	675	365	98	23.2354861	5.075423887	-2.194729606	Down	1.57E-50	7.35E-50	-
CCG018415.1	2046	475	98	9.97586703	1.674443364	-2.574760655	Down	8.09E-79	4.10E-78	AAEL000471-PA OS=Aedes aegypti GN=AAEL000471 PE=4 SV=1

CCG003745.1	936	479	98	21.9898535	3.660161457	-2.586858798	Down	6.89E-80	3.50E-79	AAEL008646-PA (Fragment) OS=Aedes aegypti GN=AAEL008646 PE=4 SV=1
CCG002448.1	894	667	98	32.0590749	3.832115351	-3.064519903	Down	2.10E-132	1.17E-131	AGAP000342-PA OS=Anopheles gambiae GN=AgaP_AGAP000342 PE=3 SV=1
CCG008959.2	804	833	98	44.519638	4.261083487	-3.385149637	Down	2.45E-181	1.44E-180	Carbonic anhydrase OS=Aedes aegypti PE=2 SV=1
CCG025683.1	841	4	99	0.20437448	4.115183592	4.331669824	Up	2.95E-05	5.04E-05	-
CCG004651.1	850	7	99	0.3538684	4.071611059	3.524314902	Up	4.31E-06	7.69E-06	AAEL001997-PA OS=Aedes aegypti GN=AAEL001997 PE=4 SV=1
CCG010806.1	1587	9	99	0.2436847	2.180762067	3.161744822	Up	1.09E-06	2.01E-06	AAEL013562-PA OS=Aedes aegypti GN=AAEL013562 PE=4 SV=1
CCG010518.1	2164	9	99	0.17870962	1.599292699	3.161744822	Up	1.09E-06	2.01E-06	Uncharacterized protein OS=Danaus plexippus GN=KGM_14575 PE=4 SV=1
CCG020532.1	1632	13	99	0.34228343	2.12063076	2.631230106	Up	6.34E-08	1.22E-07	AAEL002075-PA OS=Aedes aegypti GN=AAEL002075 PE=4 SV=1
CCG021581.1	783	13	99	0.71341833	4.420012006	2.631230106	Up	6.34E-08	1.23E-07	AAEL002020-PA OS=Aedes aegypti GN=AAEL002820 PE=4 SV=1
CCG024467.1	504	19	99	1.61989079	6.866804366	2.08374231	Up	8.17E-10	1.69E-09	-
CCG010612.1	999	34	99	1.46243341	3.464333734	1.244206983	Up	4.95E-06	8.81E-06	Uncharacterized protein OS=Anopheles darlingi GN=AND_13575 PE=4 SV=1
CCG020741.1	1462	34	99	0.99929615	2.367215732	1.244206983	Up	4.95E-06	8.81E-06	AAEL006184-PA OS=Aedes aegypti GN=AAEL006184 PE=4 SV=1
CCG004841.1	711	164	99	9.91144371	4.867608158	-1.025882181	Down	1.26E-08	2.51E-08	AAEL004588-PA OS=Aedes aegypti GN=AAEL004588 PE=4 SV=1
CCG022519.1	1016	172	99	7.27440389	3.40636752	-1.094594931	Down	7.88E-10	1.63E-09	Putative gag-pol polyprotein OS=Aster yellows phytoplasma PE=4 SV=1
CCG021740.1	237	172	99	31.1847863	14.60282447	-1.094594931	Down	7.88E-10	1.63E-09	AAEL009810-PA OS=Aedes aegypti GN=AAEL009810 PE=2 SV=1
CCG005543.1	1641	177	99	4.63476114	2.109000244	-1.135935726	Down	1.30E-10	2.78E-10	AAEL001867-PA OS=Aedes aegypti GN=AAEL001867 PE=4 SV=1

CCG001289.1	816	222	99	11.6902954	4.24126152	-1.462746042	Down	1.71E-18	6.59E-18	Salivary chymotrypsin-like enzyme OS=Aedes aegypti PE=2 SV=1
CCG013771.1	3555	300	99	3.62613794	0.973521632	-1.897148867	Down	8.34E-35	3.61E-34	Putative uncharacterized protein OS=Amoebophilus asiaticus (strain 5a2) GN=Aasi_1435 PE=4 SV=1
CCG023402.1	885	542	99	26.3159279	3.910586893	-2.750479217	Down	1.76E-96	9.28E-96	AAEL010943-PA OS=Aedes aegypti GN=AAEL010943 PE=4 SV=1
CCG015182.1	1257	2	100	0.06836871	2.781088049	5.346169394	Up	9.36E-05	0.00015397	AAEL000429-PA OS=Aedes aegypti GN=AAEL000429 PE=4 SV=1
CCG013739.1	927	6	100	0.27812126	3.771119393	3.761206893	Up	8.38E-06	1.48E-05	AAEL000894-PA OS=Aedes aegypti GN=AAEL008945 PE=4 SV=1
CCG007526.1	366	6	100	0.70442188	9.551441741	3.761206893	Up	8.38E-06	1.48E-05	AAEL007344-PA OS=Aedes aegypti GN=AAEL007344 PE=4 SV=1
CCG026559.1	1407	13	100	0.39701958	2.484596786	2.645729675	Up	6.34E-08	1.23E-07	AAEL005737-PA OS=Aedes aegypti GN=AAEL005737 PE=4 SV=1
CCG026719.1	1077	15	100	0.59846427	3.245893851	2.439278798	Up	1.48E-08	2.92E-08	AAEL004309-PA OS=Aedes aegypti GN=AAEL004309 PE=4 SV=1
CCG001725.1	1905	20	100	0.45112582	1.835080146	2.024241299	Up	4.46E-10	9.31E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014911 PE=4 SV=1
CCG015555.1	1911	23	100	0.51716583	1.829318512	1.822607438	Up	1.33E-09	2.74E-09	-
CCG011021.1	888	26	100	1.25812286	3.93674288	1.645729675	Up	1.57E-08	3.11E-08	Lipase OS=Aedes aegypti GN=AAEL004933 PE=3 SV=1
CCG010317.1	837	40	100	2.05351181	4.176616102	1.024241299	Up	8.52E-05	0.00014049	AAEL004147-PA OS=Aedes aegypti GN=AAEL004147 PE=4 SV=1
CCG004822.1	1242	186	100	6.43508103	2.814676069	-1.192989418	Down	7.23E-12	1.62E-11	AAEL000387-PA OS=Aedes aegypti GN=AAEL000387 PE=4 SV=1
CCG015655.1	1509	290	100	8.25793442	2.316651874	-1.833739696	Down	3.10E-32	1.32E-31	AAEL012617-PA OS=Aedes aegypti GN=AAEL012617 PE=4 SV=1
CCG020458.1	951	322	100	14.5491636	3.675949187	-1.984747485	Down	1.78E-39	7.94E-39	AAEL011460-PA OS=Aedes aegypti GN=AAEL011460 PE=4 SV=1
CCG025367.1	1260	969	100	33.0457721	2.77446641	-3.574183462	Down	4.79E-221	2.90E-220	AAEL006607-PA OS=Aedes aegypti GN=AAEL006607 PE=4 SV=1
CCG025837.1	273	0	101	0.001	12.93328188	13.65880079	Up	4.44E-16	1.65E-15	-

CCG027905.1	2061	19	101	0.3961305	1.713142142	2.112597173	Up	7.92E-10	1.64E-09	AAEL013492-PA OS=Aedes aegypti GN=PPO5 PE=4 SV=1
CCG007302.1	478	35	101	3.14631948	7.386581494	1.23124167	Up	4.74E-06	8.44E-06	-
CCG015995.1	1170	186	101	6.83108602	3.017765773	-1.178634125	Down	1.14E-11	2.54E-11	AAEL014372-PA OS=Aedes aegypti GN=AAEL014372 PE=4 SV=1
CCG011315.1	900	208	101	9.93078311	3.923095504	-1.339915032	Down	1.99E-15	7.23E-15	AAEL002673-PA OS=Aedes aegypti GN=AAEL002673 PE=4 SV=1
CCG015824.1	1644	233	101	6.0899928	2.147680021	-1.503661458	Down	4.39E-20	1.71E-19	-
CCG017113.1	3342	286	101	3.6772424	1.056488915	-1.79934665	Down	5.00E-31	2.12E-30	AAEL000489-PA OS=Aedes aegypti GN=AAEL000489 PE=4 SV=1
CCG012113.1	513	297	101	24.8772148	6.882623692	-1.853794434	Down	1.90E-33	8.14E-33	AAEL005317-PB (Fragment) OS=Aedes aegypti GN=AAEL005317 PE=4 SV=1
CCG001935.1	863	566	101	28.1817726	4.09129311	-2.784133556	Down	5.33E-102	2.83E-101	Hydroxypyruvate isomerase OS=Aedes aegypti GN=AAEL002165 PE=3 SV=1
CCG027572.1	3246	858	101	11.357989	1.087734428	-3.384309151	Down	1.11E-186	6.54E-186	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010658 PE=4 SV=1
CCG007744.1	1461	0	102	0.001	2.440618912	11.25303133	Up	4.44E-16	1.63E-15	AAEL000058-PA OS=Aedes aegypti GN=CYP325Q2 PE=3 SV=1
CCG026725.1	516	3	102	0.24982404	6.910357036	4.789776045	Up	5.28E-05	8.85E-05	Fibrinogen and fibronectin OS=Culex quinquefasciatus GN=CpipJ_CPIJ018858 PE=4 SV=1
CCG026303.1	1416	4	102	0.12138343	2.518180954	4.374738546	Up	2.95E-05	5.03E-05	AGAP007008-PA OS=Anopheles gambiae GN=AgaP_AGAP007008 PE=4 SV=3
CCG024271.1	981	4	102	0.17520789	3.634805536	4.374738546	Up	2.95E-05	5.04E-05	AAEL008043-PA OS=Aedes aegypti GN=PNRLIKE PE=2 SV=2
CCG017213.1	987	5	102	0.21767849	3.612709454	4.052810451	Up	1.59E-05	2.76E-05	AAEL015060-PA (Fragment) OS=Aedes aegypti GN=AAEL015060 PE=4 SV=1
CCG028090.1	381	5	102	0.56390728	9.358908742	4.052810451	Up	1.59E-05	2.76E-05	AAEL005909-PA OS=Aedes aegypti GN=AAEL005909 PE=4 SV=1

CCG012305.1	680	8	102	0.50552629	5.243741516	3.374738546	Up	2.19E-06	3.96E-06	AAEL005746-PA OS=Aedes aegypti GN=CYP9AE1 PE=3 SV=1
CCG014116.1	633	8	102	0.54306142	5.633087252	3.374738546	Up	2.19E-06	3.96E-06	AAEL001246-PA OS=Aedes aegypti GN=AAEL001246 PE=3 SV=1
CCG011735.1	540	18	102	1.43232449	6.603230057	2.204813544	Up	1.63E-09	3.33E-09	AAEL0008910-PA OS=Aedes aegypti GN=AAEL008910 PE=4 SV=1
CCG016690.1	2481	18	102	0.3117514	1.437220569	2.204813544	Up	1.63E-09	3.33E-09	AAEL0008045-PA OS=Aedes aegypti GN=AAEL008045 PE=4 SV=1
CCG023501.1	1053	28	102	1.14259503	3.386271824	1.567383624	Up	3.49E-08	6.82E-08	AAEL002675-PA OS=Aedes aegypti GN=AAEL002675 PE=4 SV=1
CCG021208.1	507	30	102	2.54258785	7.033026096	1.46784795	Up	1.47E-07	2.79E-07	-
CCG025889.1	1197	31	102	1.11283356	2.978900778	1.420542235	Up	2.89E-07	5.43E-07	AAEL005468-PA OS=Aedes aegypti GN=AAEL005468 PE=4 SV=1
CCG019611.1	1062	192	102	7.76853959	3.357574605	-1.210223955	Down	1.88E-12	4.50E-12	AAEL004825-PA OS=Aedes aegypti GN=AAEL004825 PE=4 SV=1
CCG016169.1	2378	241	102	4.35479648	1.499471922	-1.53815079	Down	2.23E-21	8.80E-21	AAEL008427-PA OS=Aedes aegypti GN=AAEL008427 PE=4 SV=1
CCG014600.1	1281	248	102	8.31888695	2.783563022	-1.579457765	Down	9.11E-23	3.64E-22	AAEL001095-PA OS=Aedes aegypti GN=AAEL001095 PE=4 SV=1
CCG026669.1	948	255	102	11.5583147	3.761333577	-1.619614891	Down	3.52E-24	1.42E-23	AAEL000139-PA OS=Aedes aegypti GN=AAEL000139 PE=4 SV=1
CCG018497.1	1477	296	102	8.61140247	2.414180251	-1.83471482	Down	6.85E-33	2.93E-32	AAEL011086-PA OS=Aedes aegypti GN=AAEL011086 PE=4 SV=2
CCG018194.1	1590	324	102	8.75609686	2.242606434	-1.965111457	Down	3.31E-39	1.47E-38	AAEL002669-PA OS=Aedes aegypti GN=AAEL002669 PE=4 SV=1
CCG010182.1	1566	406	102	11.1403016	2.276975882	-2.290597371	Down	5.21E-59	2.50E-58	AAEL000467-PA OS=Aedes aegypti GN=AAEL006467 PE=3 SV=1
CCG022799.1	1788	582	102	13.9867928	1.994264111	-2.810136797	Down	7.03E-106	3.75E-105	AAEL005753-PA OS=Aedes aegypti GN=AAEL005753 PE=3 SV=1
CCG028627.1	11511	907	102	3.38576573	0.309768415	-3.450220195	Down	1.05E-200	6.29E-200	AAEL010965-PA OS=Aedes aegypti GN=AAEL010965 PE=4 SV=1
CCG011343.1	2271	7908	102	149.627768	1.570120753	-6.57435861	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_22379 PE=4 SV=1

CCG017579.1	468	6	103	0.55089403	7.693808777	3.80385123	Up	8.38E-06	1.48E-05	AAEL005379-PA OS=Aedes aegypti GN=AAEL005379 PE=4 SV=1
CCG006440.1	582	8	103	0.59064927	6.186774068	3.388813731	Up	2.19E-06	3.97E-06	-
CCG004083.1	1533	9	103	0.2522685	2.348794852	3.21888873	Up	1.09E-06	2.01E-06	AAEL005737-PA OS=Aedes aegypti GN=AAEL005737 PE=4 SV=1
CCG001464.1	453	12	103	1.13827112	7.948570657	2.80385123	Up	1.30E-07	2.49E-07	-
CCG018428.1	462	14	103	1.30211317	7.793728371	2.581458809	Up	3.07E-08	6.01E-08	AAEL006566-PA OS=Aedes aegypti GN=AAEL006566 PE=4 SV=1
CCG007586.1	1200	16	103	0.57292979	3.000585423	2.388813731	Up	7.09E-09	1.42E-08	AAEL004138-PA OS=Aedes aegypti GN=AAEL004138 PE=4 SV=1
CCG015063.1	591	19	103	1.38142971	6.092559234	2.140886218	Up	7.81E-10	1.62E-09	-
CCG012177.1	817	22	103	1.15707976	4.407224611	1.929382112	Up	2.40E-10	5.06E-10	moduler of mdg4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001643 PE=4 SV=1
CCG012228.1	1371	27	103	0.8462311	2.626332974	1.633926229	Up	1.13E-08	2.25E-08	AAEL009303-PA OS=Aedes aegypti GN=AAEL009303 PE=4 SV=1
CCG004270.1	2257	30	103	0.57115287	1.595348918	1.481923135	Up	1.04E-07	1.99E-07	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG004442.1	1887	200	103	4.5542909	1.908162431	-1.255042459	Down	1.37E-13	4.14E-13	AAEL010471-PA OS=Aedes aegypti GN=AAEL010471 PE=3 SV=1
CCG027430.1	568	233	103	17.6266693	6.339264978	-1.475372414	Down	1.45E-19	5.64E-19	Glutaredoxin OS=Aedes aegypti GN=AAEL001109 PE=3 SV=1
CCG000659.1	1893	270	103	6.12880525	1.902114373	-1.688001866	Down	5.56E-27	2.30E-26	AAEL001796-PA OS=Aedes aegypti GN=AAEL001796 PE=3 SV=1
CCG009458.1	1245	342	103	11.8037343	2.892130528	-2.029038784	Down	5.08E-43	2.30E-42	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010424 PE=3 SV=1
CCG000389.2	1662	347	103	8.97141872	2.16648767	-2.049978122	Down	3.38E-44	1.54E-43	AAEL010381-PA OS=Aedes aegypti GN=AAEL010381 PE=3 SV=1
CCG005542.1	981	491	103	21.5067683	3.670440884	-2.550765483	Down	1.48E-80	7.53E-80	AAEL001878-PA OS=Aedes aegypti GN=AAEL001878 PE=3 SV=1
CCG011040.1	1731	3880	103	96.3157541	2.080128543	-5.533027206	Down	0	0	AAEL013303-PA OS=Aedes aegypti GN=AAEL013303 PE=4 SV=1
CCG018637.1	675	2	104	0.12731773	5.386164125	5.402752922	Up	9.36E-05	0.00015366	-

CCG025955.1	723	8	104	0.47546041	5.028576465	3.402752922	Up	2.19E-06	3.96E-06	40S ribosomal protein S4 (Fragment) OS=Aedes triseriatus PE=2 SV=1
CCG028331.1	2001	8	104	0.17179304	1.816921931	3.402752922	Up	2.19E-06	3.97E-06	AAEL000033-PA OS=Aedes aegypti GN=AAEL000033 PE=4 SV=1
CCG015371.1	323	9	104	1.19729911	11.25591574	3.232827921	Up	1.09E-06	2.01E-06	-
CCG025032.1	1524	13	104	0.36653973	2.385604189	2.702313204	Up	6.34E-08	1.23E-07	AAEL003580-PA OS=Aedes aegypti GN=CYP4H28 PE=3 SV=2
CCG028139.1	1265	21	104	0.71333156	2.874040146	2.010435499	Up	2.15E-10	4.54E-10	Uncharacterized protein OS=Anopheles darlingi GN=AND_19488 PE=4 SV=1
CCG025533.1	655	23	104	1.50886091	5.550627152	1.879190966	Up	3.03E-10	6.36E-10	AAEL004640-PA OS=Aedes aegypti GN=AAEL004640 PE=4 SV=1
CCG011219.1	1911	24	104	0.5396513	1.902491253	1.817790421	Up	6.63E-10	1.38E-09	AAEL009923-PA (Fragment) OS=Aedes aegypti GN=AAEL009923 PE=4 SV=1
CCG020754.1	949	24	104	1.08669508	3.83104403	1.817790421	Up	6.63E-10	1.38E-09	-
CCG005380.1	2689	29	104	0.46341477	1.352049381	1.544771927	Up	3.56E-08	6.96E-08	GF20629 OS=Drosophila ananassae GN=GF20629 PE=4 SV=1
CCG013864.1	1412	38	104	1.15640929	2.574830584	1.154825409	Up	1.01E-05	1.76E-05	AAEL009143-PA OS=Aedes aegypti GN=AAEL009143 PE=4 SV=1
CCG027650.1	1113	40	104	1.54428516	3.266541585	1.080824827	Up	2.82E-05	4.82E-05	AAEL014959-PA OS=Aedes aegypti GN=AAEL014959 PE=4 SV=1
CCG000970.1	2198	229	104	4.47682858	1.65407679	-1.436450866	Down	1.49E-18	5.72E-18	AAEL011697-PA OS=Aedes aegypti GN=AAEL011697 PE=3 SV=1
CCG000958.1	952	232	104	10.471616	3.818971412	-1.455228073	Down	4.05E-19	1.56E-18	-
CCG016141.1	1191	356	104	12.8440181	3.05261191	-2.072980509	Down	6.06E-46	2.78E-45	AAEL005935-PA OS=Aedes aegypti GN=AAEL005935 PE=4 SV=1
CCG009963.1	1436	379	104	11.3408979	2.531797204	-2.163301116	Down	1.77E-51	8.31E-51	Membrane glycoprotein LIG-1 (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG014357.1	1752	431	104	10.5707509	2.075148849	-2.348791137	Down	1.91E-64	9.31E-64	AAEL014765-PA OS=Aedes aegypti GN=AAEL014765 PE=4 SV=1

CCG024981.1	1191	558	104	20.131916	3.05261191	-2.72136839	Down	3.89E-98	2.05E-97	AAEL013452-PA OS=Aedes aegypti GN=AAEL013452 PE=4 SV=1
CCG003655.1	591	0	105	0.001	6.210861355	12.60057765	Up	4.44E-16	1.66E-15	Angiopoietin-like protein 2 (Fragment) OS=Aedes albopictus PE=2 SV=1
CCG003332.1	1017	0	105	0.001	3.609261614	11.817488	Up	4.44E-16	1.63E-15	AAEL009198-PA OS=Aedes aegypti GN=AAEL009198 PE=4 SV=1
CCG016674.1	507	7	105	0.5932705	7.239879805	3.609203799	Up	4.31E-06	7.70E-06	AAEL000463-PA OS=Aedes aegypti GN=AAEL000463 PE=4 SV=1
CCG014883.1	596	12	105	0.86516244	6.158756814	2.831596221	Up	1.30E-07	2.49E-07	-
CCG011862.1	1368	26	105	0.81667624	2.683201068	1.716119003	Up	2.40E-09	4.90E-09	AAEL001541-PA OS=Aedes aegypti GN=AAEL001541 PE=4 SV=1
CCG012949.1	1911	184	105	4.13732662	1.920784438	-1.107003235	Down	1.39E-10	2.96E-10	AAEL006583-PA OS=Aedes aegypti GN=AAEL006583 PE=4 SV=1
CCG024147.1	1855	219	105	5.07297675	1.978770383	-1.358228338	Down	1.79E-16	6.72E-16	AAEL011677-PA OS=Aedes aegypti GN=AAEL011677 PE=4 SV=1
CCG026869.2	3795	235	105	2.66083996	0.967225049	-1.459958225	Down	1.96E-19	7.61E-19	AAEL011178-PA OS=Aedes aegypti GN=AAEL011178 PE=4 SV=1
CCG018352.1	201	240	105	51.3071458	18.26178637	-1.490331874	Down	2.18E-20	8.53E-20	AGAP011555-PA OS=Anopheles gambiae GN=AGAP011555 PE=4 SV=3
CCG008965.1	930	291	105	13.4453686	3.946902216	-1.768316621	Down	8.05E-31	3.41E-30	AAEL009324-PA OS=Aedes aegypti GN=AAEL009324 PE=3 SV=1
CCG012614.1	705	325	105	19.8087429	5.20655186	-1.927737186	Down	2.33E-38	1.03E-37	AAEL000168-PA OS=Aedes aegypti GN=AAEL000168 PE=4 SV=1
CCG006126.1	342	1232	105	154.791559	10.73280427	-3.850227819	Down	2.23E-298	1.41E-297	-
CCG017807.1	426	13585	105	1370.29072	8.616476669	-7.313168224	Down	0	0	AAEL004249-PA OS=Aedes aegypti GN=AAEL004249 PE=4 SV=1
CCG017814.1	474	4	106	0.36261379	7.817673709	4.430233658	Up	2.95E-05	5.04E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017096 PE=4 SV=1
CCG016672.1	351	5	106	0.61210448	10.55720039	4.108305563	Up	1.59E-05	2.76E-05	AAEL000456-PA OS=Aedes aegypti GN=AAEL000456 PE=4 SV=1
CCG012572.1	946	5	106	0.22711276	3.91710078	4.108305563	Up	1.59E-05	2.76E-05	AAEL005399-PA OS=Aedes aegypti GN=AAEL005399 PE=4 SV=1

CCG019188.1	2019	11	106	0.2341095	1.835352817	2.97080204	Up	2.67E-07	5.04E-07	Proaccrosin OS=Culex quinquefasciatus GN=CpipJ_CPIJ012481 PE=3 SV=1
CCG005420.1	1140	12	106	0.452313	3.250506437	2.845271158	Up	1.30E-07	2.49E-07	AAEL010472-PA OS=Aedes aegypti GN=AAEL010472 PE=4 SV=1
CCG003608.1	839	17	106	0.87066208	4.416659521	2.342770817	Up	3.39E-09	6.88E-09	Late trypsin OS=Aedes albopictus PE=3 SV=1
CCG019215.1	956	19	106	0.854001	3.876126922	2.182306145	Up	7.75E-10	1.61E-09	AAEL011782-PA OS=Aedes aegypti GN=AAEL011782 PE=4 SV=1
CCG013077.1	2636	20	106	0.32602227	1.405757715	2.108305563	Up	3.75E-10	7.83E-10	STIP OS=Aedes aegypti PE=2 SV=1
CCG014481.1	1929	21	106	0.46778871	1.920983586	2.037916236	Up	1.93E-10	4.08E-10	Heat shock 70 Ca OS=Aedes aegypti GN=Hsp70Ca PE=3 SV=1
CCG025256.1	1304	22	106	0.72494951	2.841700412	1.97080204	Up	1.29E-10	2.76E-10	AAEL007464-PA OS=Aedes aegypti GN=AAEL007464 PE=4 SV=1
CCG008435.1	270	27	106	4.29697346	13.72436051	1.675346156	Up	3.70E-09	7.49E-09	AAEL008777-PA OS=Aedes aegypti GN=AAEL008777 PE=4 SV=1
CCG025403.1	4422	184	106	1.78797629	0.837986734	-1.093328298	Down	2.12E-10	4.48E-10	AAEL003498-PA OS=Aedes aegypti GN=AAEL003498 PE=4 SV=1
CCG027351.1	861	221	106	11.0293976	4.303806432	-1.357668901	Down	1.33E-16	5.02E-16	AAEL003977-PA OS=Aedes aegypti GN=AAEL003977 PE=4 SV=1
CCG024891.1	1570	222	106	6.07597521	2.360240343	-1.364182208	Down	8.81E-17	3.32E-16	-
CCG011600.1	3028	228	106	3.23550181	1.223770587	-1.402656356	Down	7.07E-18	2.70E-17	-
CCG019190.1	1353	318	106	10.0993168	2.738785911	-1.882649297	Down	1.99E-36	8.69E-36	AAEL003765-PA (Fragment) OS=Aedes aegypti GN=AAEL003765 PE=4 SV=1
CCG008902.1	1361	385	106	12.1552886	2.722687243	-2.158480977	Down	4.02E-52	1.89E-51	AAEL000820-PA OS=Aedes aegypti GN=AAEL000820 PE=4 SV=1
CCG019315.1	1596	514	106	13.8386238	2.321790312	-2.575390891	Down	3.61E-85	1.85E-84	Lysine 6-oxidase OS=Culex quinquefasciatus GN=CpipJ_CPIJ006993 PE=4 SV=1
CCG019482.1	4722	963	106	8.76320509	0.784747424	-3.481158329	Down	1.11E-214	6.68E-214	AAEL000580-PA OS=Aedes aegypti GN=AAEL000580 PE=4 SV=1
CCG003726.1	1197	3	107	0.10769357	3.124925326	4.858817689	Up	5.28E-05	8.83E-05	AAEL007809-PA OS=Aedes aegypti GN=AAEL007809 PE=4 SV=1

CCG016389.1	1147	12	107	0.44955259	3.261147005	2.858817689	Up	1.30E-07	2.49E-07	-
CCG025540.1	522	16	107	1.31707999	7.16577704	2.44378019	Up	7.09E-09	1.42E-08	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001647 PE=4 SV=1
CCG001361.1	2025	36	107	0.76390639	1.847178081	1.273855189	Up	1.31E-06	2.40E-06	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG013285.1	905	38	107	1.80425405	4.133188524	1.195852677	Up	4.12E-06	7.37E-06	-
CCG005435.1	470	175	107	15.9993693	7.958586414	-1.007430922	Down	6.97E-09	1.40E-08	-
CCG010382.1	1299	177	107	5.85499848	2.879550127	-1.02382536	Down	3.57E-09	7.23E-09	AAEL009835-PA OS=Aedes aegypti GN=AAEL009835 PE=4 SV=1
CCG000440.1	1524	180	107	5.07516551	2.454419695	-1.048072906	Down	1.29E-09	2.65E-09	AAEL002640-PA OS=Aedes aegypti GN=AAEL002640 PE=4 SV=1
CCG001900.1	1233	210	107	7.31844628	3.03368663	-1.270465327	Down	1.93E-14	6.78E-14	AAEL007224-PA OS=Aedes aegypti GN=AAEL007224 PE=4 SV=1
CCG020590.1	4202	276	107	2.82238143	0.890179823	-1.664744267	Down	4.94E-27	2.04E-26	5-hydroxytryptamine 7 receptor OS=Aedes aegypti PE=2 SV=1
CCG028521.2	429	303	107	30.3492531	8.719197237	-1.799393793	Down	8.92E-33	3.81E-32	AAEL014082-PA OS=Aedes aegypti GN=AAEL014082 PE=4 SV=1
CCG010038.1	1569	323	107	8.84590458	2.384025248	-1.891610164	Down	3.32E-37	1.46E-36	AAEL001737-PA (Fragment) OS=Aedes aegypti GN=AAEL001737 PE=4 SV=1
CCG003730.1	1698	355	107	8.98366065	2.202906722	-2.027895024	Down	1.47E-44	6.68E-44	AAEL012509-PA (Fragment) OS=Aedes aegypti GN=AAEL012509 PE=4 SV=1
CCG017847.1	884	374	107	18.1795031	4.23137513	-2.10311427	Down	4.59E-49	2.13E-48	AAEL011439-PA OS=Aedes aegypti GN=AAEL011439 PE=4 SV=1
CCG003107.1	1443	416	107	12.3876712	2.592193773	-2.256659528	Down	2.44E-59	1.17E-58	AAEL002127-PA OS=Aedes aegypti GN=AAEL002127 PE=4 SV=1
CCG016269.1	2037	1221	107	25.7565272	1.836296325	-3.810067295	Down	2.90E-293	1.82E-292	AAEL007126-PA OS=Aedes aegypti GN=AAEL007126 PE=4 SV=1
CCG014506.1	561	0	108	0.001	6.729935635	12.71637699	Up	4.44E-16	1.64E-15	AAEL007209-PA OS=Aedes aegypti GN=AAEL007209 PE=3 SV=1

CCG001064.1	822	1	108	0.05227462	4.593058262	6.457200706	Up	0.000587	0.00091854	Uncharacterized protein OS=Tetrapisispora phaffii (strain ATCC 24235 / CBS 4417 / NBRC 1672 / NRRL Y-8282 / UCD 70-5) GN=TPHA0B00750 PE=4 SV=1
CCG007760.1	1269	7	108	0.23702769	2.975172491	3.649845784	Up	4.31E-06	7.70E-06	AAEL010770-FA OS=Aedes aegypti GN=AAEL010770 PE=4 SV=1
CCG014282.1	483	16	108	1.42342806	7.816757539	2.457200706	Up	7.09E-09	1.42E-08	-
CCG006574.1	1335	21	108	0.67592841	2.828085312	2.064883283	Up	1.83E-10	3.87E-10	AAEL005730-FA OS=Aedes aegypti GN=AAEL005730 PE=4 SV=1
CCG028603.1	1524	22	108	0.62029801	2.477358196	1.997769087	Up	1.04E-10	2.22E-10	AAEL001982-FA OS=Aedes aegypti GN=AAEL001982 PE=4 SV=1
CCG004183.1	747	23	108	1.32303065	5.05420869	1.93363875	Up	9.22E-11	1.97E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001628 PE=4 SV=1
CCG001920.1	1198	23	108	0.82496152	3.151497405	1.93363875	Up	9.22E-11	1.97E-10	AAEL001545-FA OS=Aedes aegypti GN=AAEL001545 PE=4 SV=1
CCG007108.1	1691	26	108	0.6606819	2.23269893	1.756760988	Up	7.63E-10	1.58E-09	AAEL005202-FA OS=Aedes aegypti GN=AAEL005202 PE=4 SV=1
CCG006612.1	1126	28	108	1.06851916	3.353014113	1.649845784	Up	3.88E-09	7.85E-09	AAEL005403-FA OS=Aedes aegypti GN=AAEL005403 PE=4 SV=1
CCG006191.1	1838	32	108	0.7481129	2.054131606	1.457200706	Up	7.35E-08	1.42E-07	-
CCG023031.1	543	32	108	2.53228639	6.953027424	1.457200706	Up	7.35E-08	1.42E-07	AAEL003709-FA OS=Aedes aegypti GN=AAEL003709 PE=4 SV=1
CCG012544.1	255	35	108	5.89780671	14.8058584	1.327917689	Up	5.20E-07	9.71E-07	AAEL002812-FA OS=Aedes aegypti GN=AAEL002812 PE=4 SV=1
CCG011980.1	1590	39	108	1.05397462	2.37452446	1.171798487	Up	5.28E-06	9.38E-06	AAEL004630-FA OS=Aedes aegypti GN=AAEL004630 PE=4 SV=1
CCG026948.1	723	41	108	2.4367346	5.221983252	1.099648701	Up	1.50E-05	2.61E-05	AAEL001155-FA OS=Aedes aegypti GN=AAEL001155 PE=4 SV=1
CCG028538.7	1551	184	108	5.09763454	2.434232038	-1.06636125	Down	4.85E-10	1.01E-09	AAEL005762-FA OS=Aedes aegypti GN=AAEL005762 PE=4 SV=1
CCG006688.1	3324	197	108	2.54664191	1.135828487	-1.164851113	Down	4.53E-12	1.04E-11	AAEL006394-FA OS=Aedes aegypti GN=AAEL006394 PE=4 SV=1

CCG002113.1	1726	204	108	5.07869401	2.187424039	-1.215224636	Down	3.24E-13	8.86E-13	AAEL005921-PA OS=Aedes aegypti GN=AAEL005921 PE=4 SV_1
CCG008467.1	2766	216	108	3.35555411	1.364965254	-1.297686796	Down	2.92E-15	1.06E-14	AAEL002141-PA OS=Aedes aegypti GN=AAEL002141 PE=4 SV_1
CCG017367.1	957	262	108	11.763919	3.945134683	-1.576222296	Down	6.77E-24	2.72E-23	AAEL007997-PA OS=Aedes aegypti GN=AAEL007997 PE=4 SV_1
CCG025839.1	717	357	108	21.3949725	5.265681857	-2.022579558	Down	1.19E-44	5.41E-44	AAEL014204-PA OS=Aedes aegypti GN=AAEL014204 PE=4 SV_1
CCG004657.1	1376	453	108	14.1462862	2.743818235	-2.366166534	Down	3.05E-68	1.50E-67	AAEL003183-PA OS=Aedes aegypti GN=AAEL003183 PE=3 SV_1
CCG026898.1	2046	493	108	10.3538999	1.845304932	-2.48824313	Down	1.30E-78	6.59E-78	AAEL001457-PA OS=Aedes aegypti GN=AAEL001457 PE=3 SV_1
CCG009041.1	952	1131	108	51.049128	3.965854928	-3.686182508	Down	1.58E-264	9.80E-264	AAEL002889-PA OS=Aedes aegypti GN=AAEL002889 PE=4 SV_1
CCG019749.1	1128	1390	108	52.9502935	3.347069053	-3.983668462	Down	0	0	AAEL014045-PA OS=Aedes aegypti GN=AAEL014045 PE=4 SV_1
CCG011404.1	1164	1921	108	70.9148283	3.243551453	-4.450441098	Down	0	0	AAEL015119-PA OS=Aedes aegypti GN=AAEL015119 PE=4 SV_1
CCG003581.1	1773	0	109	0.001	2.149155199	11.06955395	Up	4.44E-16	1.65E-15	-
CCG014188.1	330	12	109	1.5625358	11.54682475	2.885535028	Up	1.30E-07	2.49E-07	AAEL014451-PA OS=Aedes aegypti GN=AAEL014451 PE=4 SV_1
CCG004092.1	609	13	109	0.91725213	6.256900112	2.77005781	Up	6.34E-08	1.23E-07	AAEL004317-PA OS=Aedes aegypti GN=AAEL004317 PE=4 SV_1
CCG015750.1	1014	13	109	0.55089403	3.757842375	2.77005781	Up	6.34E-08	1.23E-07	AAEL013683-PA OS=Aedes aegypti GN=AAEL013683 PE=4 SV_1
CCG021033.1	475	16	109	1.44740159	8.022004565	2.470497529	Up	7.09E-09	1.42E-08	-
CCG011645.1	594	18	109	1.30211317	6.41490264	2.300572527	Up	1.62E-09	3.32E-09	AAEL004854-PA OS=Aedes aegypti GN=AAEL004854 PE=4 SV_1
CCG000939.1	1084	22	109	0.87207948	3.515177277	2.01106591	Up	9.70E-11	2.07E-10	AAEL008576-PA OS=Aedes aegypti GN=AAEL008576 PE=4 SV_1
CCG019555.1	4513	189	109	1.79953021	0.844327979	-1.091744896	Down	1.27E-10	2.71E-10	AAEL005920-PA OS=Aedes aegypti GN=AAEL005920 PE=4 SV_1
CCG021276.1	702	195	109	11.9360374	5.427994542	-1.136832785	Down	1.47E-11	3.26E-11	AAEL010661-PA OS=Aedes aegypti GN=AAEL010661 PE=4 SV_1

CCG018801.1	1299	259	109	8.56748365	2.933373494	-1.546310759	Down	5.02E-23	2.01E-22	AGAP002971-PA OS=Anopheles gambiae GN=AGAP002971 PE=2 SV=3
CCG000081.1	897	371	109	17.7723206	4.247995728	-2.064777848	Down	1.44E-47	6.63E-47	AAEL001799-PA OS=Aedes aegypti GN=AAEL001799 PE=4 SV=1
CCG002546.1	1143	403	109	15.1503089	3.333728931	-2.1841385	Down	2.78E-55	1.32E-54	AAEL000791-PA OS=Aedes aegypti GN=AAEL006791 PE=4 SV=1
CCG023578.1	1203	551	109	19.6810671	3.167458161	-2.63541098	Down	1.47E-93	7.66E-93	AAEL005781-PA OS=Aedes aegypti GN=AAEL005781 PE=4 SV=1
CCG022774.1	1140	905	109	34.1119384	3.342501902	-3.351276453	Down	4.78E-195	2.85E-194	AAEL013461-PA OS=Aedes aegypti GN=AAEL013461 PE=4 SV=1
CCG005216.1	1317	3	110	0.09788094	2.919825699	4.898710417	Up	5.28E-05	8.85E-05	AGAP000437-PA OS=Anopheles gambiae GN=AgaP_AGAP000437 PE=4 SV=5
CCG005923.1	900	7	110	0.33420905	4.272678272	3.676317995	Up	4.31E-06	7.69E-06	AAEL004050-PA OS=Aedes aegypti GN=AAEL004050 PE=4 SV=1
CCG018122.1	477	8	110	0.72066641	8.061657117	3.483672917	Up	2.19E-06	3.96E-06	AAEL004515-PA OS=Aedes aegypti GN=AAEL004515 PE=4 SV=1
CCG023588.1	1500	12	110	0.34375788	2.563606963	2.898710417	Up	1.30E-07	2.49E-07	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG025805.1	374	17	110	1.95316976	10.28184611	2.396210076	Up	3.39E-09	6.88E-09	Mod(Mdg4)-v12 OS=Anopheles gambiae GN=mod(mdg4) PE=4 SV=1
CCG002468.1	735	20	110	1.16924448	5.231850946	2.161744822	Up	3.70E-10	7.72E-10	AAEL008761-PA OS=Aedes aegypti GN=AAEL008761 PE=4 SV=1
CCG013717.1	1377	44	110	1.37303437	2.792600178	1.024241299	Up	3.72E-05	6.29E-05	AAEL014550-PA (Fragment) OS=Aedes aegypti GN=AAEL014550 PE=3 SV=1
CCG005383.1	1130	44	110	1.67315781	3.403018093	1.024241299	Up	3.72E-05	6.29E-05	-
CCG007747.1	1203	230	110	8.21532748	3.196517411	-1.361817134	Down	2.74E-17	1.04E-16	AAEL000044-PA OS=Aedes aegypti GN=CYP325Q1 PE=3 SV=1
CCG013773.1	1451	325	110	9.62450982	2.650179493	-1.860622991	Down	1.27E-36	5.55E-36	AAEL011086-PA OS=Aedes aegypti GN=AAEL011086 PE=4 SV=2
CCG024616.1	1050	374	110	15.3054102	3.662295662	-2.063221543	Down	6.79E-48	3.14E-47	AAEL009937-PA OS=Aedes aegypti GN=AAEL009937 PE=4 SV=1

CCG008842.1	3861	438	110	4.87457751	0.995962301	-2.291114142	Down	1.53E-63	7.45E-63	AAEL004824-PA (Fragment) OS=Aedes aegypti GN=AAEL004824 PE=4 SV=1
CCG024108.1	474	604	110	54.7546829	8.112680264	-2.754731822	Down	1.72E-107	9.19E-107	AAEL004120-PA OS=Aedes aegypti GN=AAEL004120 PE=4 SV=1
CCG008923.1	450	5	111	0.4774415	8.623041604	4.174800975	Up	1.59E-05	2.76E-05	AAEL000583-PA OS=Aedes aegypti GN=AAEL000583 PE=3 SV=1
CCG015606.1	558	14	111	1.0780937	6.95406581	2.689374148	Up	3.07E-08	6.01E-08	AGAP007888-PA OS=Anopheles gambiae GN=AGAP007888 PE=4 SV=4
CCG026760.2	801	16	111	0.85832179	4.844405395	2.49672907	Up	7.09E-09	1.42E-08	AAEL011985-PA OS=Aedes aegypti GN=AAEL011985 PE=4 SV=1
CCG011274.1	730	21	111	1.23611565	5.315573591	2.104411647	Up	1.78E-10	3.76E-10	-
CCG003251.1	1329	30	111	0.96997144	2.91976578	1.589838475	Up	6.03E-09	1.21E-08	AAEL006438-PA OS=Aedes aegypti GN=AAEL006438 PE=4 SV=1
CCG000068.1	1260	42	111	1.43232449	3.079657716	1.104411647	Up	1.06E-05	1.86E-05	AAEL015181-PA OS=Aedes aegypti GN=AAEL015181 PE=4 SV=1
CCG007001.1	1527	188	111	5.29031441	2.541171396	-1.057859782	Down	4.10E-10	8.55E-10	AAEL003966-PA OS=Aedes aegypti GN=AAEL003966 PE=4 SV=1
CCG014555.1	2922	203	111	2.98523481	1.32798382	-1.168606847	Down	1.89E-12	4.52E-12	AAEL006199-PA OS=Aedes aegypti GN=AAEL006199 PE=4 SV=1
CCG001840.1	1074	231	111	9.24209376	3.613006259	-1.355019971	Down	3.09E-17	1.17E-16	AAEL008164-PA OS=Aedes aegypti GN=AAEL008164 PE=4 SV=1
CCG022565.1	483	297	111	26.4223834	8.033889693	-1.717590051	Down	3.21E-30	1.35E-29	AAEL000316-PA OS=Aedes aegypti GN=AAEL000316 PE=4 SV=1
CCG003565.1	861	389	111	19.4137361	4.506816169	-2.106897275	Down	4.43E-51	2.08E-50	AAEL001301-PA OS=Aedes aegypti GN=AAEL001301 PE=4 SV=1
CCG019102.1	2451	767	111	13.4466693	1.583177773	-3.086353697	Down	4.67E-153	2.66E-152	Microtubule-associated protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013260 PE=4 SV=1
CCG015978.1	705	2	112	0.12189996	5.553655317	5.509668126	Up	9.36E-05	0.00015369	AAEL003906-PA OS=Aedes aegypti GN=AAEL003906 PE=4 SV=1
CCG023600.1	714	6	112	0.36109021	5.483651258	3.924705625	Up	8.38E-06	1.47E-05	-
CCG007485.1	2278	8	112	0.15090337	1.718756365	3.509668126	Up	2.19E-06	3.96E-06	AAEL011567-PA OS=Aedes aegypti GN=AAEL011567 PE=3 SV=1

CCG009569.1	996	19	112	0.81970377	3.931051203	2.261740612	Up	7.73E-10	1.60E-09	AAEL007611-PA OS=Reedes aegypti GN=AAEL007611 PE=4 SV=1
CCG025031.1	1599	22	112	0.59120335	2.448609755	2.050236507	Up	8.76E-11	1.87E-10	AAEL003366-PA OS=Reedes aegypti GN=AAEL003366 PE=4 SV=1
CCG026571.1	639	27	112	1.81562259	6.127272298	1.754780624	Up	3.80E-10	7.93E-10	AAEL013522-PA OS=Reedes aegypti GN=AAEL013522 PE=4 SV=1
CCG018408.1	984	28	112	1.22271603	3.978990852	1.702313204	Up	8.67E-10	1.79E-09	AAEL004507-PA OS=Reedes aegypti GN=AAEL004507 PE=4 SV=1
CCG027442.1	1668	28	112	0.72131449	2.347318344	1.702313204	Up	8.67E-10	1.79E-09	AAEL008626-PA OS=Reedes aegypti GN=AAEL008626 PE=4 SV=1
CCG025790.1	1054	39	112	1.58996172	3.714731498	1.224265907	Up	1.58E-06	2.89E-06	-
CCG026593.1	2223	44	112	0.85050307	1.761280701	1.050236507	Up	2.16E-05	3.71E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009352 PE=4 SV=1
CCG027822.1	900	399	112	19.0499157	4.350363332	-2.13057681	Down	4.36E-53	2.06E-52	AAEL008106-PA OS=Reedes aegypti GN=AAEL008106 PE=4 SV=1
CCG006063.1	357	451	112	54.2838944	10.96730252	-2.307315497	Down	6.20E-66	3.04E-65	AAEL005948-PA OS=Reedes aegypti GN=AAEL005948 PE=4 SV=1
CCG009207.1	2309	1041	112	19.3726694	1.695680814	-3.514086227	Down	7.46E-234	4.55E-233	AAEL002555-PA OS=Reedes aegypti GN=AAEL002555 PE=3 SV=1
CCG008377.1	13638	1701	112	5.35940157	0.287089529	-4.2224993	Down	0	0	AAEL003155-PA OS=Reedes aegypti GN=AAEL003155 PE=4 SV=1
CCG015526.1	1983	0	113	0.001	1.992075277	10.96005645	Up	4.44E-16	1.64E-15	AGAP003785-PB OS=Anopheles gambiae GN=AgaP_AGAP003785 PE=3 SV=1
CCG005833.1	1200	3	113	0.10742434	3.291904396	4.937529666	Up	5.28E-05	8.84E-05	-
CCG012237.1	1042	4	113	0.164951	3.791060725	4.522492166	Up	2.95E-05	5.03E-05	AAEL013813-PA OS=Reedes aegypti GN=AAEL013813 PE=4 SV=1
CCG025994.1	828	23	113	1.19360374	4.770875936	1.99893021	Up	4.67E-11	1.01E-10	AAEL005268-PA OS=Reedes aegypti GN=AAEL005268 PE=4 SV=1
CCG007030.1	885	25	113	1.21383431	4.463599181	1.878635976	Up	5.34E-11	1.15E-10	AAEL004183-PA OS=Reedes aegypti GN=AAEL004183 PE=4 SV=1
CCG010099.1	683	29	113	1.82448361	5.783726611	1.664511171	Up	1.33E-09	2.74E-09	AAEL009735-PA OS=Reedes aegypti GN=AAEL009735 PE=4 SV=1

CCG022398.1	840	35	113	1.79040561	4.702720566	1.393209149	Up	1.01E-07	1.93E-07	Transcriptional activator HATZ OS=Culex quinquefasciatus GN=CpipJ_CPIJ017497 PE=4 SV=1
CCG017176.1	585	35	113	2.57083882	6.752624402	1.393209149	Up	1.01E-07	1.93E-07	AAEL011303-PA OS=Aedes aegypti GN=AAEL011303 PE=3 SV=1
CCG017622.1	1157	38	113	1.4112791	3.414248293	1.274564653	Up	6.50E-07	1.20E-06	AAEL000170-PA OS=Aedes aegypti GN=AAEL000170 PE=4 SV=1
CCG014213.1	710	45	113	2.72343388	5.563782078	1.03063907	Up	2.65E-05	4.53E-05	AAEL003463-PA OS=Aedes aegypti GN=AAEL003463 PE=4 SV=1
CCG002925.1	1779	198	113	4.78246625	2.220508867	-1.106864454	Down	2.80E-11	6.11E-11	AAEL000575-PA OS=Aedes aegypti GN=AAEL000575 PE=3 SV=1
CCG015247.1	2271	261	113	4.9383975	1.739447501	-1.50541383	Down	2.40E-22	9.53E-22	AAEL002684-PA OS=Aedes aegypti GN=AAEL002684 PE=4 SV=1
CCG001545.1	1948	271	113	5.97782242	2.027867184	-1.559656875	Down	2.63E-24	1.06E-23	-
CCG000557.1	1719	528	113	13.1983827	2.29801354	-2.521901953	Down	2.23E-85	1.15E-84	AAEL013371-PA OS=Aedes aegypti GN=AAEL013371 PE=4 SV=1
CCG025274.1	2338	786	113	14.4457705	1.689600203	-3.095893336	Down	2.78E-157	1.59E-156	AAEL013884-PA OS=Aedes aegypti GN=AAEL013884 PE=4 SV=1
CCG008526.1	483	1	114	0.08896425	8.251021847	6.535203218	Up	0.000587	0.00091621	Uncharacterized protein OS=Anopheles darlingi GN=AND_17523 PE=4 SV=1
CCG006107.1	643	1	114	0.06682696	6.197890439	6.535203218	Up	0.000587	0.00091711	-
CCG021862.1	1212	3	114	0.10636073	3.288154746	4.950240717	Up	5.28E-05	8.85E-05	AAEL005945-PA (Fragment) OS=Aedes aegypti GN=GPRDOP4 PE=3 SV=2
CCG027300.1	855	3	114	0.150771	4.66110357	4.950240717	Up	5.28E-05	8.85E-05	AAEL004145-PA OS=Aedes aegypti GN=AAEL004145 PE=4 SV=1
CCG005681.1	363	6	114	0.71024355	10.97863237	3.950240717	Up	8.38E-06	1.48E-05	AAEL001341-PA OS=Aedes aegypti GN=AAEL001341 PE=4 SV=1
CCG026051.1	456	9	114	0.84808687	8.739569193	3.365278217	Up	1.09E-06	2.01E-06	-
CCG001290.1	477	12	114	1.08099961	8.354808285	2.950240717	Up	1.30E-07	2.49E-07	AAEL014061-PA (Fragment) OS=Aedes aegypti GN=AAEL014061 PE=4 SV=1
CCG018260.1	660	21	114	1.36721883	6.038247806	2.142885795	Up	1.76E-10	3.73E-10	AAEL009202-PA OS=Aedes aegypti GN=AAEL009202 PE=4 SV=1

CCG024529.1	1925	21	114	0.46876074	2.070256391	2.142885795	Up	1.76E-10	3.73E-10	Eukaryotic translation initiation factor 4e type OS=Culex quinquefasciatus GN=CpipJ_CPIJ012021 PE=2 SV=2
CCG019843.1	1740	22	114	0.5432955	2.290369857	2.075771599	Up	8.54E-11	1.83E-10	AGAP009563-PA (Fragment) OS=Anopheles gambiae GN=AGAP009563 PE=4 SV=3
CCG022236.1	942	34	114	1.5509246	4.230619482	1.447740377	Up	3.69E-08	7.20E-08	AAEL013581-PA OS=Aedes aegypti GN=AAEL013581 PE=4 SV=1
CCG016756.1	1865	190	114	4.37761371	2.136859813	-1.03465239	Down	6.84E-10	1.42E-09	AAEL010478-PA OS=Aedes aegypti GN=AAEL010478 PE=3 SV=1
CCG028109.1	2085	258	114	5.31711824	1.911387795	-1.476024037	Down	1.65E-21	6.50E-21	AAEL003115-PA OS=Aedes aegypti GN=AAEL003115 PE=4 SV=1
CCG017815.1	1080	371	114	14.7608996	3.690040326	-2.000072159	Down	1.12E-45	5.12E-45	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017097 PE=4 SV=1
CCG021742.1	894	414	114	19.8987362	4.457766837	-2.15828374	Down	6.51E-56	3.10E-55	AAEL008108-PA OS=Aedes aegypti GN=AAEL008108 PE=4 SV=1
CCG001012.1	1741	519	114	12.8094729	2.289054309	-2.48438751	Down	1.62E-82	8.27E-82	AAEL001241-PA OS=Aedes aegypti GN=AAEL001241 PE=4 SV=1
CCG001893.1	1416	624	114	18.9358153	2.814437537	-2.750199001	Down	8.57E-111	4.61E-110	AAEL011177-PA OS=Aedes aegypti GN=AAEL011177 PE=4 SV=1
CCG003439.1	1416	882	114	26.7650466	2.814437537	-3.249431628	Down	9.84E-185	5.79E-184	AAEL011177-PA OS=Aedes aegypti GN=AAEL011177 PE=4 SV=1
CCG028190.2	471	1155	114	105.371642	8.461238964	-3.638473918	Down	3.70E-267	2.30E-266	AAEL008289-PA OS=Aedes aegypti GN=AAEL008289 PE=4 SV=1
CCG019655.1	1536	2	115	0.05595018	2.617318899	5.547803255	Up	9.36E-05	0.00015359	AAEL002743-PA OS=Aedes aegypti GN=AAEL002743 PE=4 SV=1
CCG028515.2	462	7	115	0.65105659	8.70173556	3.740448333	Up	4.31E-06	7.71E-06	-
CCG001402.1	981	8	115	0.35041578	4.098065065	3.547803255	Up	2.19E-06	3.97E-06	AAEL003007-PA OS=Aedes aegypti GN=AAEL003007 PE=4 SV=1
CCG011498.1	708	10	115	0.60691716	5.678251171	3.22587516	Up	5.43E-07	1.01E-06	Pipsqueak OS=Culex quinquefasciatus GN=CpipJ_CPIJ013751 PE=4 SV=1
CCG010987.1	1224	11	115	0.38616592	3.284478618	3.088371636	Up	2.67E-07	5.03E-07	-

CCG014107.1	2661	12	115	0.19377558	1.510786106	2.962840754	Up	1.30E-07	2.49E-07	AAEL014653-PA OS=Aedes aegypti GN=AAEL014653 PE=4 SV=2
CCG018162.1	1356	13	115	0.41195173	2.964750611	2.847363537	Up	6.34E-08	1.23E-07	AAEL001732-PA OS=Aedes aegypti GN=AAEL001732 PE=4 SV=1
CCG000937.1	276	17	115	2.64668655	14.56594866	2.460340414	Up	3.39E-09	6.89E-09	AAEL0008578-PA OS=Aedes aegypti GN=AAEL008578 PE=4 SV=1
CCG018854.1	828	22	115	1.14170792	4.855316218	2.088371636	Up	8.49E-11	1.82E-10	AAEL000487-PA OS=Aedes aegypti GN=AAEL000487 PE=4 SV=1
CCG022953.1	2004	36	115	0.7719114	2.006088737	1.377878253	Up	9.91E-08	1.90E-07	AAEL004950-PA OS=Aedes aegypti GN=AAEL004950 PE=4 SV=1
CCG027039.1	633	44	115	2.98683779	6.351029745	1.088371636	Up	9.37E-06	1.64E-05	AAEL0006340-PA OS=Aedes aegypti GN=AAEL006340 PE=4 SV=1
CCG022137.1	816	235	115	12.3748623	4.926717927	-1.328713692	Down	4.96E-17	1.88E-16	AGAP001434-PA OS=Anopheles gambiae GN=AGAP001434 PE=3 SV=3
CCG019018.1	1581	235	115	6.3870257	2.542822156	-1.328713692	Down	4.96E-17	1.88E-16	AAEL013602-PA (Fragment) OS=Aedes aegypti GN=AAEL013602 PE=4 SV=1
CCG027114.1	4964	338	115	2.92581996	0.80987144	-1.853076182	Down	8.08E-38	3.56E-37	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010720 PE=4 SV=1
CCG020414.1	1578	500	115	13.6152518	2.547656419	-2.41798103	Down	4.34E-77	2.19E-76	AAEL0006846-PA OS=Aedes aegypti GN=AAEL006846 PE=4 SV=1
CCG016989.1	816	544	115	28.6464897	4.926717927	-2.539659586	Down	1.28E-88	6.64E-88	Uncharacterized protein OS=Anopheles darlingi GN=AND_10976 PE=4 SV=1
CCG012984.1	933	3866	115	178.050369	4.308897994	-5.368822667	Down	0	0	AAEL013766-PA OS=Aedes aegypti GN=AAEL013766 PE=4 SV=1
CCG018515.1	849	6	116	0.30367304	4.776395884	3.975331698	Up	8.38E-06	1.47E-05	AAEL004333-PA OS=Aedes aegypti GN=AAEL004333 PE=4 SV=1
CCG013884.1	1091	13	116	0.51201334	3.716920353	2.859854481	Up	6.34E-08	1.23E-07	AAEL002098-PA (Fragment) OS=Aedes aegypti GN=AAEL002098 PE=4 SV=1

CCG023702.1	1497	18	116	0.51667016	2.708857786	2.390369198	Up	1.62E-09	3.32E-09	AAEL015342-PA OS=Aedes aegypti GN=AAEL015342 PE=4 SV=1
CCG024657.1	1155	24	116	0.8928776	3.51096113	1.975331698	Up	2.44E-11	5.32E-11	AAEL012784-PA OS=Aedes aegypti GN=AAEL012784 PE=4 SV=1
CCG018075.1	1194	27	116	0.97167742	3.396281495	1.805406697	Up	8.21E-11	1.76E-10	AAEL002541-PA OS=Aedes aegypti GN=AAEL002541 PE=4 SV=1
CCG005239.1	657	33	116	2.15829717	6.172237604	1.51590008	Up	9.17E-09	1.83E-08	AAEL011755-PA OS=Aedes aegypti GN=AAEL011755 PE=3 SV=1
CCG017330.1	1284	215	116	7.19508796	3.158224381	-1.187898651	Down	2.08E-13	5.97E-13	AAEL003606-PA OS=Aedes aegypti GN=AAEL003606 PE=4 SV=1
CCG002876.1	1419	223	116	6.75281946	2.85775906	-1.240605701	Down	9.77E-15	3.49E-14	AAEL015239-PA (Fragment) OS=Aedes aegypti GN=AAEL015239 PE=4 SV=1
CCG027210.1	858	256	116	12.8208066	4.726293829	-1.439705801	Down	1.26E-20	4.96E-20	-
CCG009424.1	720	270	116	16.1136505	5.632166813	-1.516521398	Down	2.67E-23	1.07E-22	AAEL008136-PA OS=Aedes aegypti GN=AAEL008136 PE=4 SV=1
CCG021109.1	543	287	116	22.7114435	7.468066493	-1.604612728	Down	1.13E-26	4.67E-26	AAEL013327-PA OS=Aedes aegypti GN=AAEL013327 PE=4 SV=1
CCG026084.1	2442	288	116	5.06768369	1.660589724	-1.609630802	Down	7.11E-27	2.94E-26	AAEL004047-PA (Fragment) OS=Aedes aegypti GN=AAEL004047 PE=4 SV=1
CCG022848.1	960	300	116	13.4280421	4.22412511	-1.668524492	Down	2.47E-29	1.04E-28	AAEL005485-PA OS=Aedes aegypti GN=AAEL005485 PE=4 SV=1
CCG017359.1	2385	365	116	6.57608098	1.700276774	-1.951458455	Down	1.46E-43	6.63E-43	AAEL007186-PA OS=Aedes aegypti GN=AAEL007186 PE=4 SV=2
CCG024024.1	1914	2979	116	66.8792264	2.118683441	-4.980318209	Down	0	0	-
CCG006350.1	528	7	117	0.56967451	7.74643633	3.765323001	Up	4.31E-06	7.69E-06	AAEL002875-PA OS=Aedes aegypti GN=AAEL002875 PE=4 SV=1
CCG021510.1	807	9	117	0.47921637	5.06830035	3.402752922	Up	1.09E-06	2.01E-06	AAEL001326-PA OS=Aedes aegypti GN=AAEL001326 PE=4 SV=1
CCG015752.1	525	12	117	0.98216536	7.790701681	2.987715423	Up	1.30E-07	2.49E-07	AAEL013681-PA OS=Aedes aegypti GN=AAEL013681 PE=4 SV=1
CCG024664.1	1620	13	117	0.34481886	2.524764434	2.872238205	Up	6.34E-08	1.23E-07	AAEL004399-PA (Fragment) OS=Aedes aegypti GN=GPRFSH PE=3 SV=2

CCG007365.1	1182	14	117	0.50894779	3.460337041	2.765323001	Up	3.07E-08	6.01E-08	AAEL007981-PA OS=Aedes aegypti GN=AAEL007981 PE=4 SV=1
CCG013632.1	925	17	117	0.78971404	4.421749603	2.485215082	Up	3.39E-09	6.88E-09	AAEL004924-PA OS=Aedes aegypti GN=AAEL004924 PE=4 SV=1
CCG021240.1	1164	26	117	0.95980507	3.513847408	1.872238205	Up	2.67E-11	5.81E-11	AAEL005701-PA OS=Aedes aegypti GN=AAEL005701 PE=4 SV=1
CCG021837.1	1104	28	117	1.08981211	3.704817375	1.765323001	Up	1.29E-10	2.75E-10	AAEL006132-PA OS=Aedes aegypti GN=AAEL006132 PE=4 SV=1
CCG027040.1	1673	29	117	0.74484298	2.444780862	1.714696928	Up	2.96E-10	6.21E-10	AAEL006356-PA OS=Aedes aegypti GN=AAEL006356 PE=4 SV=1
CCG003995.1	408	33	117	3.47549324	10.02479996	1.528283804	Up	6.44E-09	1.29E-08	AAEL005390-PA (Fragment) OS=Aedes aegypti GN=AAEL005390 PE=4 SV=1
CCG006345.1	2175	220	117	4.34636396	1.880514199	-1.20868179	Down	4.98E-14	1.68E-13	AAEL002294-PA (Fragment) OS=Aedes aegypti GN=AAEL002294 PE=4 SV=1
CCG011830.1	1544	299	117	8.32121156	2.649040403	-1.651323751	Down	7.84E-29	3.29E-28	AAEL010478-PA OS=Aedes aegypti GN=AAEL010478 PE=3 SV=1
CCG017007.1	894	340	117	16.3419572	4.57507649	-1.836713013	Down	1.37E-37	6.04E-37	AAEL008899-PA OS=Aedes aegypti GN=AAEL008899 PE=3 SV=1
CCG013875.1	1396	399	117	12.2814643	2.929884228	-2.067567013	Down	4.05E-51	1.90E-50	AAEL010864-PA OS=Aedes aegypti GN=AAEL010864 PE=4 SV=1
CCG021129.1	873	436	117	21.4602569	4.685129877	-2.195506401	Down	4.73E-60	2.28E-59	AAEL002018-PA OS=Aedes aegypti GN=AAEL002018 PE=4 SV=1
CCG020220.1	1818	521	117	12.3142089	2.249790089	-2.452461639	Down	1.36E-81	6.92E-81	AAEL001374-PA OS=Aedes aegypti GN=AAEL001374 PE=4 SV=1
CCG012225.1	519	576	117	47.688954	7.880767596	-2.597247078	Down	3.85E-96	2.03E-95	AAEL009300-PA OS=Aedes aegypti GN=AAEL009300 PE=4 SV=1
CCG015062.1	1281	5	118	0.16771949	3.220200358	4.263028158	Up	1.59E-05	2.76E-05	AAEL008050-PA OS=Aedes aegypti GN=AAEL008050 PE=4 SV=1
CCG024668.1	530	5	118	0.40537485	7.783163508	4.263028158	Up	1.59E-05	2.76E-05	-
CCG019007.1	1269	8	118	0.27088879	3.250651426	3.584956253	Up	2.19E-06	3.96E-06	AAEL010089-PA OS=Aedes aegypti GN=AAEL010089 PE=4 SV=1
CCG019416.1	1019	10	118	0.42168532	4.048161589	3.263028158	Up	5.43E-07	1.01E-06	AAEL012839-PA OS=Aedes aegypti GN=AAEL012839 PE=4 SV=1

CCG015435.1	1164	11	118	0.40607138	3.543880291	3.125524635	Up	2.67E-07	5.03E-07	AAEL007757-PA OS=Aedes aegypti GN=AAEL007757 PE=4 SV=1
CCG023722.1	1056	13	118	0.52898348	3.906322594	2.884516535	Up	6.34E-08	1.23E-07	AAEL002641-PA OS=Aedes aegypti GN=AAEL002641 PE=4 SV=1
CCG014046.1	606	14	118	0.99270014	6.807057193	2.777601331	Up	3.07E-08	6.01E-08	-
CCG004662.1	1329	19	118	0.61431524	3.103895154	2.33702874	Up	7.73E-10	1.60E-09	AAEL009129-PA OS=Aedes aegypti GN=CYP6Z9 PE=3 SV=1
CCG000501.1	420	24	118	2.45541341	9.821611093	1.999993752	Up	2.15E-11	4.70E-11	Insulin-like peptide 2 OS=Aedes aegypti PE=2 SV=2
CCG015017.1	781	24	118	1.32045279	5.281788296	1.999993752	Up	2.15E-11	4.70E-11	AAEL010667-PA OS=Aedes aegypti GN=AAEL010667 PE=3 SV=1
CCG008713.2	1062	29	118	1.17337317	3.884252975	1.726975258	Up	2.02E-10	4.28E-10	AAEL002945-PA OS=Aedes aegypti GN=AAEL002945 PE=4 SV=2
CCG020228.1	234	29	118	5.32530899	17.62853273	1.726975258	Up	2.02E-10	4.28E-10	AAEL000435-PA OS=Aedes aegypti GN=AAEL000435 PE=4 SV=1
CCG015253.1	1938	30	118	0.66516617	2.128522528	1.678065658	Up	4.56E-10	9.51E-10	AAEL000811-PA (Fragment) OS=Aedes aegypti GN=GPRMTH7 PE=4 SV=2
CCG028041.1	279	33	118	5.08244173	14.785221	1.540562134	Up	4.52E-09	9.12E-09	Uncharacterized protein OS=Anopheles darlingi GN=AND_30410 PE=4 SV=1
CCG008666.1	1269	34	118	1.15127737	3.250651426	1.497493412	Up	9.26E-09	1.85E-08	AAEL008491-PA OS=Aedes aegypti GN=AAEL008491 PE=4 SV=1
CCG012183.1	1209	37	118	1.31503737	3.411974077	1.375502888	Up	6.99E-08	1.35E-07	AAEL004635-PA OS=Aedes aegypti GN=AAEL004635 PE=4 SV=1
CCG006872.1	1749	197	118	4.83993008	2.358534396	-1.037095566	Down	3.08E-10	6.47E-10	AAEL013727-PA OS=Aedes aegypti GN=AAEL013727 PE=4 SV=1
CCG027953.2	1263	264	118	8.98179726	3.26609395	-1.459437866	Down	1.25E-21	4.94E-21	AAEL004502-PA OS=Aedes aegypti GN=AAEL004502 PE=4 SV=1
CCG009530.1	3171	324	118	4.3904743	1.300875641	-1.75489375	Down	8.77E-34	3.77E-33	AAEL006482-PA OS=Aedes aegypti GN=AAEL006482 PE=4 SV=1
CCG017268.1	1539	405	118	11.3078249	2.680361702	-2.076821845	Down	3.72E-52	1.75E-51	AAEL005169-PA OS=Aedes aegypti GN=AAEL005169 PE=3 SV=1

CCG016446.1	4833	479	118	4.25874258	0.853523	-2.318925593	Down	2.42E-70	1.20E-69	Cadherin OS=Culex quinquefasciatus GN=CpipJ_CPIJ008474 PE=4 SV=1
CCG023061.1	406	543	118	57.4693741	10.16028734	-2.499852135	Down	7.17E-87	3.70E-86	AAEL003854-PA OS=Aedes aegypti GN=AAEL003854 PE=4 SV=1
CCG005407.1	519	632	118	52.3253801	7.948124584	-2.718824495	Down	8.46E-111	4.55E-110	AAEL001429-PA OS=Aedes aegypti GN=AAEL001429 PE=4 SV=1
CCG021686.1	2790	1362	118	20.9766231	1.4785221	-3.826554735	Down	0	0	AAEL002506-PA OS=Aedes aegypti GN=AAEL002506 PE=4 SV=1
CCG000646.1	870	1	119	0.0493905	4.781649352	6.597130967	Up	0.000587	0.00091626	Putative secreted protein OS=Aedes aegypti PE=2 SV=1
CCG017324.1	634	4	119	0.27110243	6.5615693	4.597130967	Up	2.95E-05	5.03E-05	-
CCG017853.1	429	5	119	0.50081276	9.697051133	4.275202872	Up	1.59E-05	2.76E-05	AAEL004909-PA OS=Aedes aegypti GN=AAEL004909 PE=4 SV=1
CCG011970.1	909	6	119	0.28362861	4.576496079	4.012168466	Up	8.38E-06	1.48E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000030 PE=4 SV=1
CCG005039.1	357	11	119	1.32399742	11.65275892	3.137699348	Up	2.67E-07	5.04E-07	-
CCG025696.1	471	11	119	1.00353945	8.832345936	3.137699348	Up	2.67E-07	5.03E-07	AAEL006313-PA OS=Aedes aegypti GN=AAEL006313 PE=4 SV=1
CCG024632.1	790	12	119	0.65270483	5.265867007	3.012168466	Up	1.30E-07	2.49E-07	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG003362.1	842	16	119	0.81652702	4.940659069	2.597130967	Up	7.09E-09	1.42E-08	AAEL007171-PA OS=Aedes aegypti GN=AAEL007171 PE=3 SV=1
CCG025750.1	267	17	119	2.73590071	15.58065519	2.509668126	Up	3.39E-09	6.89E-09	Uncharacterized protein OS=Anopheles darlingi GN=AND_13112 PE=4 SV=1
CCG007528.1	900	20	119	0.95488299	4.62226104	2.275202872	Up	3.69E-10	7.70E-10	AAEL011172-PA OS=Aedes aegypti GN=AAEL011172 PE=4 SV=1
CCG007083.1	717	21	119	1.25852779	5.802001305	2.204813544	Up	1.76E-10	3.72E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016896 PE=4 SV=1
CCG017913.1	1458	31	119	0.91362262	2.853247555	1.642934657	Up	6.94E-10	1.44E-09	AAEL010868-PA OS=Aedes aegypti GN=AAEL010868 PE=3 SV=1

CCG015401.1	543	39	119	3.08622403	7.661206143	1.311728748	Up	1.77E-07	3.35E-07	AAEL004301-PA OS=Aedes aegypti GN=AAEL004301 PE=2 SV=1
CCG018396.1	1282	42	119	1.40774482	3.244957048	1.204813544	Up	1.02E-06	1.87E-06	AAEL001811-PA OS=Aedes aegypti GN=AAEL001811 PE=4 SV=1
CCG012175.1	1321	45	119	1.4637684	3.149155894	1.105277871	Up	5.00E-06	8.89E-06	Mueller or mag4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001651 PE=4 SV=1
CCG018346.1	933	199	119	9.1650345	4.458772707	-1.039493653	Down	2.32E-10	4.89E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006398 PE=3 SV=1
CCG011374.1	4983	546	119	4.70830325	0.834845462	-2.495626174	Down	3.60E-87	1.86E-86	AAEL006004-PA OS=Aedes aegypti GN=AAEL006004 PE=4 SV=1
CCG027919.1	4169	662	119	6.82321044	0.997849589	-2.77355644	Down	1.65E-118	9.02E-118	AAEL004488-PA OS=Aedes aegypti GN=AAEL004488 PE=3 SV=1
CCG014881.1	273	715	119	112.539781	15.23822321	-2.884668465	Down	2.66E-133	1.48E-132	Putative antennal carrier protein AP-2 OS=Anopheles gambiae PE=4 SV=1
CCG025609.1	782	771	119	42.365301	5.31973777	-2.993456083	Down	3.40E-149	1.93E-148	AAEL000297-PA OS=Aedes aegypti GN=AAEL000297 PE=4 SV=1
CCG014303.1	708	0	120	0.001	5.925131656	12.5326315	Up	4.44E-16	1.63E-15	-
CCG020081.1	1698	4	120	0.10122435	2.470549595	4.609203799	Up	2.95E-05	5.02E-05	-
CCG022080.1	589	8	120	0.58362967	7.122229563	3.609203799	Up	2.19E-06	3.97E-06	AAEL000550-PA OS=Aedes aegypti GN=AAEL000550 PE=4 SV=1
CCG002337.1	867	11	120	0.54517541	4.838515816	3.149772181	Up	2.67E-07	5.04E-07	AAEL006009-PA OS=Aedes aegypti GN=AAEL006009 PE=4 SV=1
CCG005879.1	1107	13	120	0.50461296	3.789515097	2.908764081	Up	6.34E-08	1.23E-07	AAEL002330-PA OS=Aedes aegypti GN=AAEL002330 PE=4 SV=1
CCG018217.1	330	21	120	2.73443766	12.71210064	2.216886377	Up	1.76E-10	3.72E-10	AAEL001177-PA OS=Aedes aegypti GN=AAEL001177 PE=4 SV=1
CCG026304.1	309	22	120	3.05933386	13.57602981	2.149772181	Up	8.39E-11	1.80E-10	AAEL003171-PA OS=Aedes aegypti GN=AAEL003171 PE=4 SV=1
CCG008564.1	793	26	120	1.40884376	5.290029272	1.908764081	Up	1.12E-11	2.49E-11	-
CCG023091.1	691	26	120	1.61680622	6.0709019	1.908764081	Up	1.12E-11	2.49E-11	AAEL001358-PA OS=Aedes aegypti GN=AAEL001358 PE=4 SV=1

CCG024073.1	1427	30	120	0.90335812	2.939728951	1.702313204	Up	2.15E-10	4.54E-10	Flap endonuclease 1 OS=Anopheles darlingi GN=Fen1 PE=3 SV=1
CCG023881.1	2276	243	120	4.58771771	1.843142888	-1.315608704	Down	2.59E-17	9.81E-17	AAEL001794-PB (Fragment) OS=Aedes aegypti GN=AAEL001794 PE=4 SV=1
CCG012662.1	1300	328	120	10.8415946	3.226917856	-1.748348205	Down	5.22E-34	2.25E-33	AAEL000049-PA OS=Aedes aegypti GN=AAEL000049 PE=4 SV=1
CCG015146.1	2415	343	120	6.10294781	1.737057231	-1.812860967	Down	2.97E-37	1.30E-36	AAEL000309-PA OS=Aedes aegypti GN=AAEL003099 PE=3 SV=1
CCG017240.1	924	362	120	16.8344631	4.540035944	-1.890642088	Down	1.81E-41	8.13E-41	AAEL001209-PA OS=Aedes aegypti GN=AAEL001209 PE=4 SV=1
CCG018156.1	2973	382	120	5.52117007	1.411030344	-1.968225029	Down	5.10E-46	2.34E-45	AAEL000304-PA OS=Aedes aegypti GN=AAEL003043 PE=4 SV=1
CCG022219.1	1089	533	120	21.0311006	3.85215171	-2.448787923	Down	2.75E-83	1.41E-82	AAEL011623-PA OS=Aedes aegypti GN=AAEL011623 PE=4 SV=1
CCG022147.1	930	656	120	30.3098343	4.51074539	-2.748348205	Down	2.69E-116	1.46E-115	AAEL000315-PA OS=Aedes aegypti GN=AAEL003154 PE=4 SV=1
CCG027249.1	664	712	120	46.0759805	6.317760862	-2.866529632	Down	7.12E-132	3.96E-131	AAEL000447-PA OS=Aedes aegypti GN=AAEL004470 PE=4 SV=1
CCG006103.1	552	5	121	0.38921861	7.662955597	4.299248346	Up	1.59E-05	2.76E-05	-
CCG026932.1	596	10	121	0.7209687	7.097234043	3.299248346	Up	5.43E-07	1.01E-06	-
CCG025164.1	528	11	121	0.8952028	8.01127176	3.161744822	Up	2.67E-07	5.03E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003878 PE=4 SV=1
CCG017217.3	2459	18	121	0.31454055	1.72019174	2.45125144	Up	1.62E-09	3.32E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007691 PE=3 SV=1
CCG022156.1	1584	26	121	0.7053113	2.67042392	1.920736723	Up	9.00E-12	2.01E-11	AAEL013549-PA OS=Aedes aegypti GN=AAEL013549 PE=4 SV=1
CCG013799.1	383	29	121	3.25358304	11.04425976	1.763195446	Up	6.44E-11	1.39E-10	AAEL000317-PA OS=Aedes aegypti GN=AAEL003176 PE=4 SV=1
CCG024536.1	774	35	121	1.94307585	5.465053604	1.491893424	Up	6.58E-09	1.32E-08	AAEL000549-PA OS=Aedes aegypti GN=AAEL005499 PE=3 SV=1

CCG022720.1	508	36	121	3.0450993	8.32667616	1.45125144	Up	1.32E-08	2.61E-08	AAEL014169-PA OS=Aedes aegypti GN=AAEL014169 PE=4 SV=1
CCG025036.1	675	40	121	2.54635464	6.266594799	1.299248346	Up	1.71E-07	3.25E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003437 PE=4 SV=1
CCG023530.2	2063	227	121	4.72812882	2.050388507	-1.205372046	Down	2.26E-14	7.92E-14	AAEL001646-PA OS=Aedes aegypti GN=AAEL001646 PE=3 SV=1
CCG021349.1	1443	322	121	9.58853399	2.931359314	-1.709740437	Down	2.00E-32	8.55E-32	AAEL003387-PA (Fragment) OS=Aedes aegypti GN=AAEL003387 PE=4 SV=1
CCG025670.1	1437	661	121	19.7654799	2.94359881	-2.74733002	Down	4.01E-117	2.19E-116	AAEL000993-PA OS=Aedes aegypti GN=AAEL000993 PE=3 SV=1
CCG024163.1	696	702	121	43.3401634	6.077516508	-2.834150779	Down	1.70E-128	9.42E-128	AAEL006777-PA OS=Aedes aegypti GN=AAEL006777 PE=4 SV=1
CCG019834.1	2637	996	121	16.2297519	1.604077167	-3.338825491	Down	9.79E-214	5.89E-213	AAEL010028-PA OS=Aedes aegypti GN=AAEL010028 PE=3 SV=1
CCG023921.1	798	1512	121	81.4163393	5.30069109	-3.941065983	Down	0	0	AAEL012639-PA OS=Aedes aegypti GN=AAEL012639 PE=4 SV=1
CCG005992.1	636	0	122	0.001	6.705832966	12.71120083	Up	4.44E-16	1.65E-15	-
CCG021205.1	735	5	122	0.29231112	5.802598321	4.311122446	Up	1.59E-05	2.76E-05	-
CCG009690.1	960	9	122	0.40284126	4.44261434	3.46312554	Up	1.09E-06	2.01E-06	AAEL014658-PA OS=Aedes aegypti GN=AAEL014658 PE=3 SV=1
CCG018153.1	952	10	122	0.45136276	4.479947233	3.311122446	Up	5.43E-07	1.01E-06	AAEL008095-PA OS=Aedes aegypti GN=AAEL008095 PE=4 SV=1
CCG004171.1	699	13	122	0.799151	6.101444587	2.932610823	Up	6.34E-08	1.23E-07	AAEL007932-PA OS=Aedes aegypti GN=AAEL007932 PE=4 SV=1
CCG009802.1	636	13	122	0.87831219	6.705832966	2.932610823	Up	6.34E-08	1.23E-07	AGAP005722-PA OS=Anopheles gambiae GN=AGAP005722 PE=4 SV=1
CCG019881.1	1941	20	122	0.44275873	2.19727448	2.311122446	Up	3.69E-10	7.70E-10	SuHAKinin receptor OS=Culex quinquefasciatus GN=CpipJ_CPIJ005574 PE=3 SV=1
CCG003208.1	1265	24	122	0.81523607	3.371470171	2.048088041	Up	1.97E-11	4.32E-11	AAEL001631-PA OS=Aedes aegypti GN=AAEL001631 PE=4 SV=1
CCG022528.1	934	34	122	1.56420875	4.566284546	1.5455877	Up	2.26E-09	4.62E-09	-

CCG010301.1	1068	35	122	1.40818419	3.993361204	1.503767524	Up	4.64E-09	9.37E-09	AAEL012506-PA OS=Aedes aegypti GN=AAEL012506 PE=4 SV=1
CCG002388.1	1848	35	122	0.81382073	2.307851605	1.503767524	Up	4.64E-09	9.37E-09	AAEL012037-PA OS=Aedes aegypti GN=AAEL012037 PE=4 SV=1
CCG008452.2	1293	39	122	1.29607088	3.298460763	1.347648323	Up	6.71E-08	1.29E-07	AAEL011622-PA OS=Aedes aegypti GN=AAEL011622 PE=4 SV=1
CCG013890.1	1374	45	122	1.40730572	3.104010019	1.141197445	Up	2.11E-06	3.84E-06	3-oxoacyl-[acyl-carrier-protein] synthase OS=Aedes aegypti GN=AAEL002113 PE=3 SV=1
CCG026367.1	1128	200	122	7.61874727	3.780948374	-1.010805648	Down	5.33E-10	1.11E-09	AAEL014422-PA OS=Aedes aegypti GN=AAEL014422 PE=4 SV=1
CCG004240.1	1709	325	122	8.17154111	2.49555867	-1.711245367	Down	9.46E-33	4.04E-32	AAEL000989-PA OS=Aedes aegypti GN=CYP6AG7 PE=3 SV=1
CCG019263.1	1788	353	122	8.48339839	2.38529629	-1.830473832	Down	8.68E-39	3.85E-38	AAEL007793-PA OS=Aedes aegypti GN=AAEL007793 PE=4 SV=1
CCG003128.1	1485	671	122	19.4159542	2.871993109	-2.757118415	Down	2.51E-119	1.37E-118	AAEL007090-PA OS=Aedes aegypti GN=AAEL007090 PE=4 SV=1
CCG016909.1	1870	752	122	17.2798077	2.28070041	-2.92153831	Down	5.67E-142	3.20E-141	AAEL005112-PA (Fragment) OS=Aedes aegypti GN=AAEL005112 PE=4 SV=2
CCG026734.1	288	3	123	0.4476014	14.93009737	5.059865208	Up	5.28E-05	8.84E-05	AAEL003843-PA OS=Aedes aegypti GN=AAEL003843 PE=4 SV=1
CCG016077.1	1092	5	123	0.19674787	3.937608098	4.322899614	Up	1.59E-05	2.76E-05	AAEL000081-PA (Fragment) OS=Aedes aegypti GN=AAEL000081 PE=3 SV=1
CCG002190.1	763	13	123	0.73211868	5.63547581	2.944387991	Up	6.34E-08	1.23E-07	AAEL012512-PA OS=Aedes aegypti GN=IAP9 PE=4 SV=1
CCG017774.1	1437	14	123	0.41863346	2.992253335	2.837472787	Up	3.07E-08	6.01E-08	AAEL012916-PA OS=Aedes aegypti GN=AAEL012916 PE=4 SV=1
CCG004570.1	1185	16	123	0.58018207	3.628580627	2.644827709	Up	7.09E-09	1.42E-08	AAEL013003-PA OS=Aedes aegypti GN=AAEL013003 PE=4 SV=1
CCG020613.1	716	24	123	1.4403263	6.005402295	2.059865208	Up	1.96E-11	4.29E-11	-
CCG021160.1	1398	24	123	0.73767785	3.075728214	2.059865208	Up	1.96E-11	4.28E-11	AAEL007981-PA OS=Aedes aegypti GN=AAEL007981 PE=4 SV=1

CCG012178.1	813	37	123	1.95557218	5.288890582	1.435374344	Up	1.31E-08	2.60E-08	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG008663.1	1617	37	123	0.98322831	2.65916391	1.435374344	Up	1.31E-08	2.60E-08	AAEL008472-PA OS=Aedes aegypti GN=AAEL008472 PE=4 SV=1
CCG004371.1	5003	40	123	0.34355175	0.859457934	1.322899614	Up	9.04E-08	1.74E-07	-
CCG027502.1	945	43	123	1.9552366	4.550124913	1.218562954	Up	5.29E-07	9.87E-07	AAEL010768-PA OS=Aedes aegypti GN=AAEL010768 PE=3 SV=1
CCG004878.1	1599	50	123	1.34364398	2.689098213	1.000971519	Up	1.85E-05	3.18E-05	-
CCG013953.1	1521	223	123	6.29996766	2.827000686	-1.156072191	Down	2.52E-13	7.07E-13	AAEL011854-PA OS=Aedes aegypti GN=AAEL011854 PE=4 SV=1
CCG013651.1	861	479	123	23.905346	4.994039539	-2.259054137	Down	3.77E-68	1.86E-67	AAEL003977-PA OS=Aedes aegypti GN=AAEL003977 PE=4 SV=1
CCG020078.1	2826	584	123	8.87980361	1.521538586	-2.54499685	Down	3.13E-95	1.64E-94	Ubiquitin-protein ligase OS=Culex quinquefasciatus GN=CpipJ_CPIJ002472 PE=4 SV=1
CCG014930.1	2187	683	123	13.4194462	1.966103358	-2.770914059	Down	4.38E-122	2.41E-121	AAEL010917-PA OS=Aedes aegypti GN=AAEL010917 PE=4 SV=1
CCG010808.1	846	1326	123	67.3497259	5.082586339	-3.728037351	Down	0	0	AGAP005471-PA OS=Anopheles gambiae GN=AGAP005471 PE=4 SV=4
CCG001189.1	541	5	124	0.39713248	8.012617966	4.334581419	Up	1.59E-05	2.76E-05	-
CCG024060.1	684	10	124	0.62821249	6.33746538	3.334581419	Up	5.43E-07	1.01E-06	AAEL005704-PA OS=Aedes aegypti GN=AAEL005704 PE=4 SV=1
CCG015937.1	993	11	124	0.47599907	4.365384008	3.197077896	Up	2.67E-07	5.03E-07	Putative uncharacterized protein OS=Trichoplax adhaerens GN=TRIADDRAFT_51596 PE=4 SV=1
CCG017954.1	1407	11	124	0.33593965	3.080900014	3.197077896	Up	2.67E-07	5.04E-07	AAEL007461-PA OS=Aedes aegypti GN=AAEL007461 PE=4 SV=1
CCG027703.1	2750	14	124	0.21875501	1.57630048	2.849154592	Up	3.07E-08	6.02E-08	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG004182.1	523	18	124	1.47888188	8.288386845	2.486584513	Up	1.62E-09	3.33E-09	AAEL000561-PA OS=Aedes aegypti GN=AAEL000561 PE=4 SV=1

CCG008202.1	1390	19	124	0.58735608	3.118580086	2.408582001	Up	7.73E-10	1.60E-09	AAEL015598-PA OS=Aedes aegypti GN=AAEL015598 PE=4 SV=1
CCG010871.1	1599	20	124	0.53745759	2.7109608	2.334581419	Up	3.69E-10	7.70E-10	AAEL002349-PA OS=Aedes aegypti GN=AAEL002349 PE=4 SV=1
CCG018145.1	458	23	124	2.15786877	9.464686288	2.132947558	Up	4.01E-11	8.67E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011907 PE=4 SV=1
CCG004278.1	519	39	124	3.2289396	8.352266512	1.371107295	Up	3.49E-08	6.81E-08	AAEL014250-PA OS=Aedes aegypti GN=AAEL014250 PE=4 SV=1
CCG028548.1	1860	40	124	0.92408031	2.330551785	1.334581419	Up	6.55E-08	1.26E-07	AAEL009176-PA OS=Aedes aegypti GN=GNBPB3 PE=4 SV=1
CCG022271.1	2136	41	124	0.8247936	2.029413071	1.29895751	Up	1.21E-07	2.31E-07	AAEL003211-PA OS=Aedes aegypti GN=AAEL003211 PE=4 SV=1
CCG028602.1	1239	42	124	1.45660117	3.498649168	1.264192091	Up	2.19E-07	4.14E-07	AAEL001976-PA OS=Aedes aegypti GN=AAEL001976 PE=4 SV=1
CCG022676.1	3831	210	124	2.3554279	1.131513004	-1.057736003	Down	3.96E-11	8.58E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013821 PE=4 SV=1
CCG018336.1	1698	238	124	6.02284855	2.552901248	-1.238308249	Down	1.38E-15	5.02E-15	AAEL003295-PA OS=Aedes aegypti GN=AAEL003295 PE=4 SV=1
CCG019410.1	1434	258	124	7.73095644	3.022891436	-1.354717741	Down	4.57E-19	1.77E-18	-
CCG018966.1	867	340	124	16.8508763	4.999799677	-1.752881422	Down	2.57E-35	1.12E-34	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003741 PE=4 SV=1
CCG016638.1	747	658	124	37.8501812	5.802980348	-2.70543426	Down	1.16E-114	6.30E-114	-
CCG027106.1	507	3	125	0.25425878	8.61890453	5.083134988	Up	5.28E-05	8.84E-05	AAEL014453-PA OS=Aedes aegypti GN=AAEL014453 PE=4 SV=1
CCG012765.1	1070	3	125	0.12047589	4.083910838	5.083134988	Up	5.28E-05	8.84E-05	-
CCG009169.1	1788	9	125	0.21629061	2.443951117	3.498172487	Up	1.09E-06	2.01E-06	AAEL007839-PA OS=Aedes aegypti GN=AAEL007839 PE=4 SV=1
CCG005959.1	724	24	125	1.42441109	6.035614084	2.083134988	Up	1.94E-11	4.25E-11	-
CCG025693.1	2223	25	125	0.48324038	1.965715068	2.024241299	Up	9.65E-12	2.15E-11	AAEL014407-PA OS=Aedes aegypti GN=AAEL014407 PE=4 SV=1
CCG023147.1	1257	25	125	0.85460888	3.476360061	2.024241299	Up	9.65E-12	2.15E-11	FXa-directed anticoagulant OS=Aedes albopictus PE=2 SV=1

CCG027049.1	698	26	125	1.60059183	6.260436385	1.96765777	Up	5.46E-12	1.24E-11	AAEL009170-PA OS=Aedes aegypti GN=AAEL009170 PE=3 SV=1
CCG007007.1	885	28	125	1.35949443	4.937609714	1.860742566	Up	6.71E-12	1.51E-11	AAEL012732-PA OS=Aedes aegypti GN=AAEL012732 PE=4 SV=1
CCG012170.1	573	28	125	2.0997427	7.626151128	1.860742566	Up	6.71E-12	1.51E-11	AAEL010510-PA OS=Aedes aegypti GN=AAEL010510 PE=4 SV=1
CCG027508.1	1574	38	125	1.03738876	2.776229096	1.420169975	Up	1.30E-08	2.59E-08	-
CCG023984.1	1466	40	125	1.17243478	2.980753476	1.346169394	Up	4.74E-08	9.21E-08	-
CCG013273.1	432	46	125	4.575481	10.11524212	1.144535532	Up	1.49E-06	2.72E-06	AAEL001777-PA OS=Aedes aegypti GN=AAEL001777 PE=4 SV=1
CCG010628.1	2526	256	125	4.35481079	1.729922643	-1.331902512	Down	1.77E-18	6.78E-18	AAEL013473-PA OS=Aedes aegypti GN=AAEL013473 PE=3 SV=1
CCG015758.1	1850	292	125	6.78225	2.362045728	-1.52172707	Down	3.44E-25	1.40E-24	AAEL000904-PA OS=Aedes aegypti GN=CYP6AG5 PE=3 SV=2
CCG000237.1	648	361	125	23.9383861	6.743494748	-1.827757538	Down	1.50E-39	6.67E-39	Carsymenini-1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000075 PE=4 SV=1
CCG023191.1	1362	403	125	12.714246	3.208358735	-1.98653854	Down	4.86E-49	2.26E-48	SLC4-like anion exchanger OS=Aedes aegypti PE=2 SV=1
CCG014456.1	1329	434	125	14.0322534	3.288024527	-2.093453744	Down	2.47E-56	1.18E-55	AAEL009199-PA OS=Aedes aegypti GN=AAEL009199 PE=4 SV=1
CCG016254.1	1842	764	125	17.8224089	2.372304341	-2.90933134	Down	1.39E-143	7.86E-143	AAEL013372-PA OS=Aedes aegypti GN=AAEL013372 PE=3 SV=1
CCG020907.1	786	1152	125	62.9785423	5.559522387	-3.501827513	Down	8.57E-258	5.29E-257	AAEL014108-PA OS=Aedes aegypti GN=AAEL014108 PE=4 SV=1
CCG011381.1	260	8239	125	1361.64478	16.80686383	-6.340156039	Down	0	0	AAEL000000-PA OS=Aedes aegypti GN=AAEL006000 PE=4 SV=1
CCG006380.1	933	12	126	0.5526654	4.721053455	3.094630627	Up	1.30E-07	2.49E-07	AAEL002496-PA OS=Aedes aegypti GN=AAEL002496 PE=4 SV=1
CCG003141.1	486	12	126	1.0609811	9.063256941	3.094630627	Up	1.30E-07	2.49E-07	AAEL013616-PA OS=Aedes aegypti GN=AAEL013616 PE=4 SV=1
CCG007057.1	1152	19	126	0.70870222	3.823561522	2.431665614	Up	7.73E-10	1.60E-09	AAEL006556-PA OS=Aedes aegypti GN=AAEL006556 PE=4 SV=1
CCG015074.1	1227	22	126	0.77044349	3.589847493	2.220161509	Up	8.38E-11	1.79E-10	AAEL009021-PA OS=Aedes aegypti GN=AAEL009021 PE=4 SV=1

CCG022548.1	1089	24	126	0.9469914	4.044759296	2.094630627	Up	1.94E-11	4.24E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000812 PE=4 SV=1
CCG005275.1	3237	27	126	0.35841299	1.360748493	1.924705625	Up	3.76E-12	8.72E-12	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG009582.1	1347	42	126	1.33981355	3.270039253	1.287275705	Up	1.17E-07	2.24E-07	AAEL004995-PA (Fragment) OS=Aedes aegypti GN=AAEL004995 PE=4 SV=1
CCG003065.1	1559	44	126	1.21274427	2.825364255	1.220161509	Up	3.73E-07	6.99E-07	Cytochrome P450 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003375 PE=3 SV=1
CCG001250.1	1485	46	126	1.33104902	2.966156817	1.156031171	Up	1.11E-06	2.04E-06	AAEL015072-PA OS=Aedes aegypti GN=AAEL015072 PE=4 SV=1
CCG018778.1	657	242	126	15.8275126	6.704327052	-1.23927011	Down	7.69E-16	2.81E-15	ETS transcription factor E74A OS=Aedes aegypti GN=E74A PE=2 SV=1
CCG015315.1	1464	403	126	11.8284174	3.008704148	-1.975042901	Down	1.13E-48	5.24E-48	AAEL014361-PA OS=Aedes aegypti GN=AAEL014361 PE=4 SV=1
CCG007024.2	1553	405	126	11.205887	2.836280021	-1.98218497	Down	3.90E-49	1.81E-48	AAEL007488-PA OS=Aedes aegypti GN=AAEL007488 PE=4 SV=1
CCG001536.1	1740	447	126	11.0387767	2.531461421	-2.124537894	Down	4.68E-59	2.25E-58	AAEL003472-PA OS=Aedes aegypti GN=AAEL003472 PE=4 SV=1
CCG022006.1	1346	548	126	17.4943645	3.272468702	-2.418438956	Down	2.44E-84	1.25E-83	AAEL007090-PA OS=Aedes aegypti GN=AAEL007090 PE=4 SV=1
CCG008223.1	2046	625	126	13.1261408	2.152855754	-2.608119252	Down	1.17E-104	6.25E-104	AGAP001797-PA OS=Anopheles gambiae GN=AgaP_AGAP001797 PE=3 SV=4
CCG025266.1	1725	1157	126	28.8208597	2.553474129	-3.496580022	Down	1.46E-258	9.03E-258	AAEL008943-PA OS=Aedes aegypti GN=AAEL008943 PE=4 SV=1
CCG021199.1	1269	0	127	0.001	3.498582467	11.77255478	Up	4.44E-16	1.65E-15	AAEL003128-PA OS=Aedes aegypti GN=AAEL003128 PE=4 SV=1
CCG028017.1	1035	12	127	0.49819982	4.289566329	3.10603539	Up	1.30E-07	2.49E-07	AAEL013201-PA OS=Aedes aegypti GN=AAEL013201 PE=4 SV=1

CCG012402.1	546	29	127	2.28227528	8.131320788	1.833016895	Up	6.80E-12	1.53E-11	AAEL007332-PA OS=Aedes aegypti GN=AAEL007332 PE=4 SV=1
CCG004532.1	297	38	127	5.49781116	14.94848872	1.443070377	Up	6.61E-09	1.33E-08	AAEL010223-PA OS=Aedes aegypti GN=AAEL010223 PE=4 SV=1
CCG015625.1	1257	46	127	1.57248034	3.531981822	1.167435935	Up	8.30E-07	1.53E-06	AAEL007864-PA OS=Aedes aegypti GN=AAEL007864 PE=4 SV=1
CCG008122.1	983	47	127	2.0545041	4.516481333	1.136409039	Up	1.41E-06	2.57E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014066 PE=4 SV=1
CCG016469.1	10080	211	127	0.89946567	0.440446543	-1.030101298	Down	9.27E-11	1.98E-10	AAEL008761-PA OS=Aedes aegypti GN=AAEL008761 PE=4 SV=1
CCG009497.1	447	293	127	28.1658439	9.932217338	-1.503758964	Down	7.44E-25	3.02E-24	AAEL001194-PA OS=Aedes aegypti GN=AAEL001194 PE=4 SV=1
CCG023794.1	3563	350	127	4.22099554	1.246057017	-1.760213221	Down	1.64E-36	7.19E-36	AAEL006476-PA OS=Aedes aegypti GN=AAEL006476 PE=4 SV=1
CCG010884.1	462	450	127	41.8536376	9.609742749	-2.122783301	Down	2.23E-59	1.07E-58	AAEL002309-PA OS=Aedes aegypti GN=TPX4 PE=4 SV=1
CCG019239.1	2685	710	127	11.3625741	1.653519981	-2.780677324	Down	2.55E-127	1.41E-126	AAEL005793-PA OS=Aedes aegypti GN=AAEL005793 PE=4 SV=1
CCG022052.1	1851	1117	127	25.9304125	2.39854195	-3.43441558	Down	1.00E-245	6.17E-245	AAEL004014-PA OS=Aedes aegypti GN=AAEL004014 PE=3 SV=1
CCG011184.1	1044	0	128	0.001	4.286072248	12.06544045	Up	4.44E-16	1.64E-15	AAEL013701-PA OS=Aedes aegypti GN=AAEL013701 PE=4 SV=1
CCG020527.1	414	2	128	0.20758326	10.8083561	5.702313204	Up	9.36E-05	0.0001533	-
CCG006381.1	1452	12	128	0.35512177	3.081721368	3.117350703	Up	1.30E-07	2.49E-07	AAEL002462-PA OS=Aedes aegypti GN=AAEL002462 PE=4 SV=1
CCG025211.1	1447	20	128	0.59391478	3.092370025	2.380385109	Up	3.69E-10	7.70E-10	LINE-1 retrotransposon-like element, putative (Fragment) OS=Aspergillus flavus (strain ATCC 200026 / FGSC A1120 / NRRL 3357 / JCM 12722 / SRRCL167) GN=AFLA_063980 PE=4 SV=1
CCG028375.1	682	23	128	1.44912595	6.561084204	2.178751248	Up	4.01E-11	8.66E-11	AAEL009373-PA OS=Aedes aegypti GN=AAEL009373 PE=4 SV=1

CCG024248.1	609	31	128	2.18729355	7.347552425	1.748116893	Up	2.36E-11	5.14E-11	AAEL015251-PA OS=Aedes aegypti GN=AAEL015251 PE=4 SV=1
CCG010319.1	1007	40	128	1.70684149	4.443554545	1.380385109	Up	1.77E-08	3.50E-08	AAEL006587-PA OS=Aedes aegypti GN=AAEL006587 PE=4 SV=1
CCG020949.1	393	43	128	4.70152313	11.38590185	1.276048449	Up	1.13E-07	2.16E-07	AAEL002486-PA OS=Aedes aegypti GN=AAEL002486 PE=2 SV=1
CCG000006.1	2220	213	128	4.12277183	2.015612354	-1.032396416	Down	6.98E-11	1.50E-10	AAEL001276-PA OS=Aedes aegypti GN=AAEL001276 PE=4 SV=1
CCG001847.1	1443	357	128	10.6307659	3.100942084	-1.77746706	Down	1.06E-37	4.70E-37	AAEL007253-PA OS=Aedes aegypti GN=AAEL007253 PE=4 SV=1
CCG025869.1	342	1258	128	158.058264	13.08379949	-3.594603003	Down	8.06E-288	5.05E-287	AGAP007501-PA OS=Anopheles gambiae GN=AGAP007501 PE=4 SV=4
CCG003550.1	1251	7	129	0.24043816	3.604810315	3.906185537	Up	4.31E-06	7.69E-06	AGAP001085-PA (Fragment) OS=Anopheles gambiae GN=AGAP001085 PE=4 SV=1
CCG007199.1	1245	10	129	0.34513843	3.622182894	3.391612364	Up	5.43E-07	1.01E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013333 PE=3 SV=1
CCG027970.1	1260	18	129	0.61385335	3.57906167	2.543615458	Up	1.62E-09	3.32E-09	AAEL013971-PA OS=Aedes aegypti GN=AAEL013971 PE=4 SV=1
CCG005303.1	705	20	129	1.21899956	6.396620856	2.391612364	Up	3.69E-10	7.71E-10	AAEL007570-PA OS=Aedes aegypti GN=AAEL007570 PE=4 SV=1
CCG024756.1	843	23	129	1.17236524	5.349487193	2.189978503	Up	4.01E-11	8.66E-11	AAEL005243-PA OS=Aedes aegypti GN=CLIPB13A PE=3 SV=1
CCG008854.1	771	28	129	1.56050917	5.849050199	1.906185537	Up	2.28E-12	5.42E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006402 PE=4 SV=1
CCG025034.1	954	33	129	1.48637447	4.727062582	1.66914634	Up	8.21E-11	1.76E-10	AAEL011524-PA OS=Aedes aegypti GN=AAEL011524 PE=4 SV=1
CCG002513.1	1069	34	129	1.3666707	4.218538544	1.626077618	Up	1.80E-10	3.81E-10	AAEL005465-PA OS=Aedes aegypti GN=AAEL005465 PE=4 SV=1
CCG023749.1	2562	35	129	0.58701823	1.760194264	1.584257442	Up	3.85E-10	8.05E-10	AAEL004320-PA OS=Aedes aegypti GN=AAEL004320 PE=4 SV=1

CCG021992.1	306	38	129	5.33611083	14.73731276	1.465612946	Up	3.33E-09	6.78E-09	AAEL012732-PA OS=Aedes aegypti GN=AAEL012732 PE=4 SV=1
CCG017441.1	659	41	129	2.67338258	6.843122464	1.355988455	Up	2.41E-08	4.75E-08	-
CCG027321.1	1491	44	129	1.26805387	3.024559157	1.254108841	Up	1.48E-07	2.82E-07	AAEL000559-PA OS=Aedes aegypti GN=AAEL000559 PE=4 SV=1
CCG024665.1	1770	47	129	1.14100425	2.547806612	1.158951608	Up	7.87E-07	1.45E-06	AAEL010856-PA OS=Aedes aegypti GN=AAEL010856 PE=4 SV=1
CCG002273.1	2342	48	129	0.88067774	1.925541291	1.128577959	Up	1.33E-06	2.43E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018868 PE=4 SV=1
CCG000960.1	1262	49	129	1.66839699	3.573389623	1.098830615	Up	2.21E-06	4.01E-06	AAEL014548-PA (Fragment) OS=Aedes aegypti GN=TPX3 PE=4 SV=2
CCG008237.1	1080	219	129	8.7133073	4.175571948	-1.0612466	Down	1.34E-11	2.96E-11	AAEL014272-PA OS=Aedes aegypti GN=AAEL014272 PE=4 SV=1
CCG015971.1	1723	304	129	7.58142735	2.61730569	-1.534387054	Down	1.83E-26	7.54E-26	AAEL007293-PA OS=Aedes aegypti GN=AAEL007293 PE=2 SV=1
CCG010039.1	1572	561	129	15.334619	2.868713552	-2.418316501	Down	2.72E-86	1.40E-85	Fatty acyl-CoA reductase 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017921 PE=4 SV=1
CCG012956.1	891	798	129	38.4846781	5.061299331	-2.926704477	Down	8.06E-151	4.58E-150	AAEL000585-PA OS=Aedes aegypti GN=AAEL006585 PE=3 SV=1
CCG016850.2	1675	0	130	0.001	2.71317969	11.40576888	Up	4.44E-16	1.66E-15	Alkaline phosphatase OS=Culex quinquefasciatus GN=CpipJ_CPIJ001263 PE=3 SV=1
CCG011546.1	324	1	130	0.13262264	14.02646908	6.724681017	Up	0.000587	0.00091706	-
CCG014279.1	1035	11	130	0.45668317	4.390894667	3.265249398	Up	2.67E-07	5.03E-07	AAEL007934-PA OS=Aedes aegypti GN=AAEL007934 PE=3 SV=1
CCG016577.1	1539	13	130	0.36296722	2.952940858	3.024241299	Up	6.34E-08	1.23E-07	AAEL010081-PA OS=Aedes aegypti GN=AAEL010081 PE=4 SV=1
CCG005796.1	498	16	130	1.38055372	9.125654579	2.724681017	Up	7.09E-09	1.42E-08	AAEL012441-PA OS=Aedes aegypti GN=AAEL012441 PE=4 SV=1
CCG000455.1	963	19	130	0.84779331	4.719185857	2.476753503	Up	7.73E-10	1.60E-09	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_13182 PE=4 SV=1

CCG006830.1	1773	22	130	0.5331834	2.563212623	2.265249398	Up	8.38E-11	1.79E-10	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG018209.1	1212	24	130	0.85088583	3.749650149	2.139718516	Up	1.93E-11	4.23E-11	AAEL006233-PA OS=Aedes aegypti GN=AAEL006233 PE=4 SV=1
CCG008497.1	1246	26	130	0.89663973	3.647332248	2.024241299	Up	4.70E-12	1.07E-11	AAEL007257-PA OS=Aedes aegypti GN=AAEL007257 PE=4 SV=1
CCG004877.1	4482	32	130	0.30678972	1.01396162	1.724681017	Up	2.52E-11	5.49E-11	-
CCG027876.1	556	38	130	2.93678042	8.173697806	1.476753503	Up	2.36E-09	4.82E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001639 PE=4 SV=1
CCG015050.1	429	43	130	4.30698972	10.5934172	1.298416262	Up	6.01E-08	1.17E-07	AAEL013776-PA OS=Aedes aegypti GN=AAEL013776 PE=4 SV=1
CCG005683.1	303	51	130	7.23252959	14.9986006	1.052255675	Up	4.48E-06	7.98E-06	AGAP007325-PA OS=Anopheles gambiae GN=AGAP007325 PE=4 SV=3
CCG008253.1	1214	241	130	8.53023562	3.743472801	-1.188208319	Down	7.40E-15	2.66E-14	AAEL007748-PA OS=Aedes aegypti GN=SCRB10 PE=4 SV=2
CCG022009.1	662	304	130	19.7323253	6.864918399	-1.523246497	Down	3.39E-26	1.39E-25	-
CCG022540.1	1125	489	130	18.6775113	4.039623094	-2.209009638	Down	1.19E-67	5.86E-67	AAEL006601-PA (Fragment) OS=Aedes aegypti GN=AAEL006601 PE=4 SV=1
CCG000288.1	279	1305	130	200.987468	16.2888028	-3.625153075	Down	1.14E-300	7.19E-300	AAEL009695-PA (Fragment) OS=Aedes aegypti GN=AAEL009695 PE=4 SV=1
CCG006519.1	576	4	131	0.29840093	7.950580308	4.735736205	Up	2.95E-05	5.02E-05	AAEL009491-PA OS=Aedes aegypti GN=AAEL009491 PE=4 SV=1
CCG021731.1	507	15	131	1.27129392	9.032611947	2.82884561	Up	1.48E-08	2.93E-08	AAEL003752-PA OS=Aedes aegypti GN=AAEL003752 PE=4 SV=1
CCG011717.1	1017	28	131	1.18304087	4.502983537	1.928381283	Up	1.66E-12	4.00E-12	AAEL005443-PA (Fragment) OS=Aedes aegypti GN=AAEL005443 PE=4 SV=1
CCG013851.1	1131	37	131	1.4057296	4.049101907	1.52628284	Up	8.22E-10	1.70E-09	AAEL001664-PA OS=Aedes aegypti GN=AAEL001664 PE=4 SV=1
CCG006213.1	1011	38	131	1.61508399	4.529707475	1.487808692	Up	1.67E-09	3.42E-09	AAEL013240-PA OS=Aedes aegypti GN=AAEL013240 PE=4 SV=1

CCG020511.1	2310	48	131	0.8928776	1.982482362	1.150773705	Up	7.45E-07	1.38E-06	AAEL014541-PA OS=Aedes aegypti GN=AAEL014541 PE=4 SV=1
CCG024365.1	680	52	131	3.28592088	6.734609202	1.035296487	Up	5.49E-06	9.75E-06	AAEL007059-PA OS=Aedes aegypti GN=AAEL007059 PE=4 SV=1
CCG013661.1	1177	288	131	10.5142596	3.890853235	-1.434188796	Down	6.97E-23	2.79E-22	-
CCG007827.1	843	387	131	19.7263194	5.432424979	-1.860453551	Down	2.75E-43	1.25E-42	AAEL005072-PA OS=Aedes aegypti GN=AAEL005072 PE=4 SV=1
CCG005974.1	873	887	131	43.6588254	5.245743708	-3.057054089	Down	4.85E-175	2.83E-174	AAEL001293-PA OS=Aedes aegypti GN=CRALBP2 PE=2 SV=1
CCG019440.1	1344	1	132	0.03197153	3.433402183	6.746707323	Up	0.000587	0.00091728	AAEL008449-PA OS=Aedes aegypti GN=AAEL008449 PE=4 SV=1
CCG015731.1	437	8	132	0.7866313	10.55947948	3.746707323	Up	2.19E-06	3.96E-06	-
CCG000951.1	1903	17	132	0.38385995	2.424851568	2.659244482	Up	3.39E-09	6.88E-09	AAEL008161-PA (Fragment) OS=Aedes aegypti GN=AAEL008161 PE=4 SV=1
CCG005883.1	639	21	132	1.4121509	7.221428066	2.3543899	Up	1.76E-10	3.73E-10	AAEL012109-PA OS=Aedes aegypti GN=AAEL012109 PE=4 SV=1
CCG012783.1	2573	22	132	0.36740543	1.79342889	2.287275705	Up	8.38E-11	1.79E-10	AAEL011355-PA OS=Aedes aegypti GN=AAEL011355 PE=3 SV=1
CCG012751.1	1218	24	132	0.84669428	3.788581719	2.161744822	Up	1.93E-11	4.23E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003370 PE=4 SV=1
CCG011181.1	705	25	132	1.52374945	6.545379481	2.102851133	Up	9.34E-12	2.08E-11	AAEL007570-PA OS=Aedes aegypti GN=AAEL007570 PE=4 SV=1
CCG007331.2	999	30	132	1.29038242	4.619111646	1.839816728	Up	2.44E-12	5.76E-12	AAEL000091-PA OS=Aedes aegypti GN=AAEL000091 PE=4 SV=1
CCG012149.1	1128	34	132	1.29518704	4.090862175	1.659244482	Up	5.94E-11	1.28E-10	AAEL012136-PA OS=Aedes aegypti GN=AAEL012136 PE=4 SV=1
CCG005448.2	1629	42	132	1.10787529	2.832714877	1.3543899	Up	1.70E-08	3.37E-08	AAEL011911-PA OS=Aedes aegypti GN=AAEL011911 PE=4 SV=1
CCG015411.1	1017	51	132	2.15482445	4.537357457	1.074281981	Up	2.58E-06	4.65E-06	AAEL004050-PA OS=Aedes aegypti GN=AAEL004050 PE=4 SV=1
CCG025817.1	1654	52	132	1.35092273	2.789898751	1.046267605	Up	4.19E-06	7.49E-06	AAEL015124-PA OS=Aedes aegypti GN=AAEL015124 PE=4 SV=1

CCG010594.1	1848	221	132	5.13869662	2.497019769	-1.041195236	Down	2.24E-11	4.90E-11	AAEL000250-PA OS=Aedes aegypti GN=AAEL000250 PE=4 SV=1
CCG007004.1	885	273	132	13.2550707	5.214115858	-1.346049818	Down	6.68E-20	2.60E-19	Seadex OS=Culex quinquefasciatus GN=CpipJ_CPIJ016520 PE=4 SV=1
CCG025370.1	2116	288	132	5.84843269	2.180762067	-1.423217678	Down	1.23E-22	4.91E-22	AAEL000621-PA OS=Aedes aegypti GN=AAEL006620 PE=4 SV=1
CCG004609.1	2537	352	132	5.96190248	1.818877625	-1.712724295	Down	2.24E-35	9.73E-35	AAEL013991-PA OS=Aedes aegypti GN=AAEL013991 PE=4 SV=1
CCG000472.1	1815	389	132	9.20949133	2.542420129	-1.856919022	Down	2.16E-43	9.78E-43	AAEL008559-PA OS=Aedes aegypti GN=AAEL008559 PE=3 SV=1
CCG022627.1	1389	446	132	13.7973374	3.322168851	-2.054192577	Down	1.81E-56	8.63E-56	AAEL001110-PA OS=Aedes aegypti GN=AAEL001110 PE=4 SV=1
CCG008575.1	807	670	132	35.6749965	5.718082446	-2.641309962	Down	1.16E-113	6.28E-113	Putative secreted protein OS=Aedes aegypti PE=2 SV=1
CCG002194.1	861	1908	132	95.2221297	5.359457066	-4.151138133	Down	0	0	AAEL004264-PA OS=Aedes aegypti GN=AAEL004264 PE=4 SV=1
CCG008173.1	3822	2	133	0.02248547	1.216496811	5.757595639	Up	9.36E-05	0.00015396	AAEL002094-PA OS=Aedes aegypti GN=AAEL002094 PE=4 SV=1
CCG002319.1	894	2	133	0.09612916	5.200727976	5.757595639	Up	9.36E-05	0.00015381	AGAP005746-PA OS=Anopheles gambiae GN=AGAP005746 PE=4 SV=3
CCG000595.1	1230	13	133	0.45415167	3.78004131	3.057155921	Up	6.34E-08	1.23E-07	AAEL000699-PA OS=Aedes aegypti GN=AAEL000699 PE=4 SV=1
CCG005340.1	444	19	133	1.83879495	10.47173606	2.509668126	Up	7.73E-10	1.60E-09	AAEL014251-PA OS=Aedes aegypti GN=IAP5 PE=4 SV=1
CCG013879.1	480	22	133	1.96944617	9.686355856	2.298164021	Up	8.38E-11	1.79E-10	AAEL009170-PA OS=Aedes aegypti GN=AAEL009170 PE=3 SV=1
CCG007998.1	615	25	133	1.74673718	7.560082619	2.11373945	Up	9.33E-12	2.08E-11	-
CCG028092.1	1188	28	133	1.01275469	3.913679134	1.950240717	Up	1.39E-12	3.36E-12	AAEL005905-PA OS=Aedes aegypti GN=AAEL005905 PE=4 SV=1
CCG003069.1	1171	28	133	1.02745736	3.970495995	1.950240717	Up	1.39E-12	3.36E-12	AAEL008821-PA OS=Aedes aegypti GN=AAEL008821 PE=4 SV=1
CCG017959.1	537	31	133	2.48056196	8.658195178	1.803399329	Up	3.60E-12	8.37E-12	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016891 PE=4 SV=1

CCG017141.1	1546	36	133	1.0005889	3.007406734	1.587670638	Up	1.93E-10	4.07E-10	AAEL005635-PA OS=Aedes aegypti GN=AAEL005635 PE=4 SV=1
CCG008665.1	1410	44	133	1.34089952	3.297482844	1.298164021	Up	4.24E-08	8.26E-08	AAEL008476-PA OS=Aedes aegypti GN=AAEL008476 PE=4 SV=1
CCG023008.1	837	46	133	2.36153858	5.554899415	1.234033683	Up	1.37E-07	2.60E-07	AAEL011338-PA OS=Aedes aegypti GN=AAEL011338 PE=4 SV=1
CCG022656.1	2388	46	133	0.82772521	1.947006202	1.234033683	Up	1.37E-07	2.60E-07	AAEL007981-PA OS=Aedes aegypti GN=AAEL007981 PE=4 SV=1
CCG009467.1	786	47	133	2.56943706	5.91533182	1.203006788	Up	2.40E-07	4.53E-07	-
CCG017633.1	1764	53	133	1.29104078	2.63574309	1.029675185	Up	5.12E-06	9.11E-06	AAEL014606-PA OS=Aedes aegypti GN=CYP9J7 PE=3 SV=1
CCG008576.1	2607	331	133	5.45568936	1.783448719	-1.613091767	Down	9.77E-31	4.13E-30	AAEL007704-PA OS=Aedes aegypti GN=AAEL007704 PE=4 SV=1
CCG022580.1	870	361	133	17.8299703	5.344196334	-1.738259388	Down	5.66E-37	2.48E-36	AAEL006921-PA OS=Aedes aegypti GN=AAEL006921 PE=4 SV=1
CCG007254.1	2868	419	133	6.27765649	1.621147424	-1.953210794	Down	8.19E-50	3.82E-49	AAEL002536-PA (Fragment) OS=Aedes aegypti GN=AAEL002536 PE=4 SV=1
CCG000813.1	600	710	133	50.8475193	7.749084685	-2.714079575	Down	5.19E-124	2.86E-123	AAEL009112-PA OS=Aedes aegypti GN=AAEL009112 PE=4 SV=1
CCG008244.1	1260	2	134	0.06820593	3.71778499	5.768402394	Up	9.36E-05	0.00015384	Putative salivary serpin OS=Aedes albopictus PE=2 SV=1
CCG025926.1	941	5	134	0.22831953	4.978118053	4.446474299	Up	1.59E-05	2.76E-05	AAEL000659-PA OS=Aedes aegypti GN=AAEL000659 PE=4 SV=1
CCG001428.1	2808	14	134	0.21423657	1.668236855	2.961047472	Up	3.07E-08	6.00E-08	AAEL003392-PA OS=Aedes aegypti GN=AAEL003392 PE=4 SV=1
CCG024893.1	1703	17	134	0.42894039	2.750680615	2.680939553	Up	3.39E-09	6.90E-09	AAEL010498-PA OS=Aedes aegypti GN=AAEL010498 PE=4 SV=1
CCG005075.1	1081	34	134	1.35149952	4.333403411	1.680939553	Up	2.82E-11	6.14E-11	AAEL006285-PA OS=Aedes aegypti GN=AAEL006285 PE=4 SV=1
CCG001996.1	1122	40	134	1.53189785	4.175052663	1.446474299	Up	2.36E-09	4.83E-09	AAEL012136-PA OS=Aedes aegypti GN=AAEL012136 PE=4 SV=1
CCG006828.1	1590	53	134	1.43232449	2.946169237	1.04048194	Up	3.91E-06	7.00E-06	AAEL002279-PA OS=Aedes aegypti GN=AAEL002279 PE=4 SV=1

CCG012195.1	870	225	134	11.1128624	5.384378261	-1.045378797	Down	1.27E-11	2.81E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_01420 PE=3 SV=1
CCG006261.1	1224	227	134	7.96906026	3.827131607	-1.058146093	Down	6.41E-12	1.44E-11	AAEL008584-PA OS=Aedes aegypti GN=AAEL008584 PE=4 SV=1
CCG001336.1	285	250	134	37.6927497	16.43652311	-1.19738189	Down	1.58E-15	5.75E-15	AAEL005523-PA OS=Aedes aegypti GN=AAEL005523 PE=4 SV=1
CCG018417.1	522	319	134	26.2592823	8.973963769	-1.549010219	Down	4.65E-28	1.93E-27	GK19301 OS=Drosophila willistoni GN=GK19301 PE=4 SV=1
CCG003947.1	2130	367	134	7.40370545	2.199253093	-1.751233859	Down	6.11E-38	2.70E-37	Serine/threonine-protein phosphatase OS=Aedes aegypti GN=AAEL003322 PE=3 SV=1
CCG011220.1	1560	405	134	11.1556042	3.002826338	-1.893375703	Down	2.78E-46	1.28E-45	AAEL014361-PA OS=Aedes aegypti GN=AAEL014361 PE=4 SV=1
CCG018926.1	535	475	134	38.150699	8.755904836	-2.123381309	Down	1.33E-62	6.45E-62	-
CCG011271.1	1648	616	134	16.0615027	2.842481242	-2.498384146	Down	3.05E-98	1.61E-97	-
CCG023249.1	945	668	134	30.3743732	4.957046653	-2.615301898	Down	4.45E-112	2.40E-111	AAEL003720-PA (Fragment) OS=Aedes aegypti GN=AAEL003720 PE=4 SV=1
CCG026879.1	571	12	135	0.90304171	8.265091706	3.1941663	Up	1.30E-07	2.49E-07	-
CCG006690.1	832	40	135	2.06585263	5.672316544	1.457200706	Up	1.68E-09	3.44E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014888 PE=4 SV=1
CCG018553.1	1294	224	135	7.43834664	3.647115428	-1.028226121	Down	2.64E-11	5.76E-11	AAEL000164-PA OS=Aedes aegypti GN=AAEL000164 PE=4 SV=1
CCG008151.1	1854	231	135	5.35383425	2.54550559	-1.072620241	Down	2.41E-12	5.70E-12	AAEL000487-PA OS=Aedes aegypti GN=AAEL006487 PE=3 SV=1
CCG008275.1	1341	305	135	9.77313129	3.519289608	-1.473536632	Down	4.41E-25	1.79E-24	AAEL002433-PA OS=Aedes aegypti GN=AAEL002433 PE=4 SV=1
CCG001833.1	7170	407	135	2.43914672	0.658210232	-1.889756183	Down	2.20E-46	1.01E-45	AAEL005720-PA OS=Aedes aegypti GN=AAEL005720 PE=4 SV=1
CCG026151.1	1836	438	135	10.2509498	2.570461527	-1.995658259	Down	1.82E-53	8.62E-53	Transporter OS=Aedes aegypti GN=AAEL003860 PE=3 SV=1
CCG016079.1	978	609	135	26.7572274	4.825529002	-2.471169617	Down	6.41E-96	3.37E-95	AAEL000049-PA OS=Aedes aegypti GN=AAEL000049 PE=4 SV=1

CCG023108.1	2381	912	135	16.458798	1.982094651	-3.053761213	Down	9.98E-180	5.84E-179	AAEL005003-PA OS=Aedes aegypti GN=AAEL005003 PE=3 SV=1
CCG011355.1	621	0	136	0.001	7.655918907	12.90235983	Up	4.44E-16	1.65E-15	-
CCG003102.1	453	12	136	1.13827112	10.49520009	3.204813544	Up	1.30E-07	2.49E-07	AAEL002157-PA OS=Aedes aegypti GN=AAEL002157 PE=4 SV=1
CCG016016.1	1425	14	136	0.4221588	3.336368871	2.982421123	Up	3.07E-08	6.00E-08	AAEL006747-PA OS=Aedes aegypti GN=AAEL006747 PE=4 SV=1
CCG019333.1	430	14	136	1.39901462	11.05657126	2.982421123	Up	3.07E-08	6.01E-08	DNA-directed RNA polymerase subunit OS=Aedes aegypti GN=AAEL010468 PE=3 SV=1
CCG006679.1	1860	22	136	0.50824417	2.556089054	2.330344426	Up	8.38E-11	1.80E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006366 PE=4 SV=1
CCG000158.1	447	22	136	2.11484152	10.63607526	2.330344426	Up	8.38E-11	1.80E-10	Transcription initiation factor IIA subunit 2 OS=Aedes aegypti GN=AAEL002140 PE=3 SV=1
CCG012176.1	459	24	136	2.24678351	10.35800793	2.204813544	Up	1.93E-11	4.23E-11	Mod(Mdg4)-h55.7b OS=Anopheles gambiae GN=mod(mdg4) PE=4 SV=1
CCG006237.1	2646	32	136	0.51966421	1.796797294	1.789776045	Up	2.66E-12	6.26E-12	Retrovirus-related Pol polyprotein from transposon TNT 1-94 OS=Trichinella spiralis GN=Tsp_01017 PE=4 SV=1
CCG022936.1	985	40	136	1.74496384	4.826726539	1.46784795	Up	1.19E-09	2.46E-09	AAEL014289-PA OS=Aedes aegypti GN=AAEL014289 PE=4 SV=1
CCG008721.1	333	1075	136	138.71611	14.27725418	-3.280344899	Down	9.10E-227	5.53E-226	AAEL006662-PA OS=Aedes aegypti GN=AAEL006662 PE=4 SV=1
CCG003570.1	867	1204	136	59.6719267	5.483651258	-3.443843632	Down	2.28E-265	1.42E-264	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014232 PE=4 SV=1
CCG020739.1	624	0	137	0.001	7.675134484	12.90597631	Up	4.44E-16	1.65E-15	AAEL008570-PA OS=Aedes aegypti GN=AAEL008570 PE=3 SV=1
CCG018706.1	1306	0	137	0.001	3.667139294	11.84043935	Up	4.44E-16	1.64E-15	AAEL007131-PA OS=Aedes aegypti GN=AAEL007131 PE=4 SV=1

CCG008860.1	882	2	137	0.09743704	5.430027118	5.800345287	Up	9.36E-05	0.00015408	Pyrroline-5-carboxylate reductase OS=Aedes aegypti GN=P5CR-1 PE=2 SV=1
CCG006977.1	906	5	137	0.23713982	5.28618534	4.478417192	Up	1.59E-05	2.76E-05	-
CCG021641.1	786	6	137	0.32801324	6.093236537	4.215382786	Up	8.38E-06	1.47E-05	AAEL012659-PA OS=Aedes aegypti GN=AAEL012659 PE=4 SV=1
CCG005937.1	2946	29	137	0.42298788	1.6256904	1.942364292	Up	7.77E-13	1.95E-12	AAEL009477-PA OS=Aedes aegypti GN=AAEL009477 PE=4 SV=1
CCG022395.1	612	53	137	3.72123519	7.825627317	1.072424832	Up	1.72E-06	3.13E-06	AAEL007994-PA OS=Aedes aegypti GN=AAEL007994 PE=4 SV=1
CCG013934.1	1743	332	137	8.18471135	2.747724566	-1.574694145	Down	8.21E-30	3.45E-29	AAEL000523-PA OS=Aedes aegypti GN=AAEL000523 PE=4 SV=1
CCG016257.1	537	366	137	29.2866348	8.918592026	-1.715354552	Down	8.56E-37	3.75E-36	Takeout OS=Aedes aegypti PE=2 SV=1
CCG016865.1	2114	1051	137	21.3629097	2.265508003	-3.237201667	Down	5.53E-219	3.34E-218	AAEL007103-PA OS=Aedes aegypti GN=AAEL007103 PE=4 SV=1
CCG012957.1	2865	3892	137	58.3728472	1.671652327	-5.125950708	Down	0	0	AAEL006584-PA OS=Aedes aegypti GN=AAEL006584 PE=4 SV=1
CCG021940.1	555	3	138	0.23226884	8.692328279	5.22587516	Up	5.28E-05	8.83E-05	AAEL002427-PA OS=Aedes aegypti GN=AAEL002427 PE=4 SV=1
CCG024063.1	1437	24	138	0.71765736	3.357162279	2.22587516	Up	1.93E-11	4.23E-11	AAEL005570-PA OS=Aedes aegypti GN=AAEL005570 PE=4 SV=1
CCG026485.1	1058	27	138	1.09658113	4.559775231	2.055950158	Up	2.29E-12	5.43E-12	AAEL009649-PA OS=Aedes aegypti GN=AAEL009649 PE=4 SV=1
CCG016686.1	2136	39	138	0.78455976	2.258540353	1.525435442	Up	2.94E-10	6.18E-10	AAEL012047-PA OS=Aedes aegypti GN=AAEL012047 PE=4 SV=1
CCG027779.1	1509	258	138	7.34671407	3.196979586	-1.200389595	Down	4.93E-16	1.80E-15	AAEL013791-PA OS=Aedes aegypti GN=AAEL013791 PE=4 SV=1
CCG001301.1	1411	315	138	9.59281814	3.419023526	-1.488370358	Down	3.22E-26	1.32E-25	AAEL004097-PA OS=Aedes aegypti GN=AAEL004097 PE=4 SV=1
CCG015773.1	504	336	138	28.6464897	9.571909116	-1.581479762	Down	2.48E-30	1.04E-29	AAEL007682-PA OS=Aedes aegypti GN=AAEL007682 PE=4 SV=1
CCG002321.1	1413	421	138	12.8027306	3.414184143	-1.906838762	Down	1.71E-48	7.92E-48	Oxysterol-binding protein OS=Aedes aegypti GN=AAEL012056 PE=3 SV=1

CCG024583.1	654	775	138	50.9197925	7.376517117	-2.78721484	Down	3.69E-139	2.07E-138	AAEL006268-PA OS=Aedes aegypti GN=AAEL006268 PE=4 SV=1
CCG007516.1	324	4901	138	649.983547	14.8896364	-5.448022771	Down	0	0	Probable GGY family antimicrobial peptide OS=Aedes aegypti PE=4 SV=1
CCG021578.1	948	3	139	0.13598017	5.125738894	5.236291776	Up	5.28E-05	8.84E-05	AAEL008256-PA OS=Aedes aegypti GN=AAEL008256 PE=3 SV=1
CCG016490.1	1056	5	139	0.20345518	4.601515598	4.499326182	Up	1.59E-05	2.76E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017323 PE=4 SV=1
CCG008019.1	744	5	139	0.2887751	6.531183429	4.499326182	Up	1.59E-05	2.76E-05	-
CCG013312.1	1050	6	139	0.24554134	4.627809973	4.236291776	Up	8.38E-06	1.47E-05	-
CCG023189.1	2167	16	139	0.31726615	2.242362931	2.821254277	Up	7.09E-09	1.42E-08	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_08943 PE=4 SV=1
CCG010959.1	462	17	139	1.58113742	10.51774994	2.733791435	Up	3.39E-09	6.88E-09	AAEL011749-PA OS=Aedes aegypti GN=AAEL011749 PE=4 SV=1
CCG027190.1	950	19	139	0.85939469	5.114947865	2.573326763	Up	7.73E-10	1.61E-09	AAEL014899-PA OS=Aedes aegypti GN=AAEL014899 PE=3 SV=1
CCG013088.1	1398	40	139	1.22946308	3.475822941	1.499326182	Up	4.22E-10	8.81E-10	AAEL004316-PA OS=Aedes aegypti GN=AAEL004316 PE=4 SV=1
CCG003197.1	1387	45	139	1.3941154	3.503388948	1.32940118	Up	1.15E-08	2.29E-08	AAEL005050-PA OS=Aedes aegypti GN=AAEL005050 PE=4 SV=1
CCG010257.1	582	46	139	3.39623332	8.349141703	1.29769232	Up	2.11E-08	4.16E-08	AAEL002910-PA OS=Aedes aegypti GN=AAEL002910 PE=4 SV=1
CCG008807.1	1419	315	139	9.53873601	3.424383701	-1.477953742	Down	5.83E-26	2.39E-25	AAEL015239-PA (Fragment) OS=Aedes aegypti GN=AAEL015239 PE=4 SV=1
CCG001337.1	1218	340	139	11.9948356	3.989491356	-1.58813666	Down	7.45E-31	3.16E-30	AAEL005531-PA OS=Aedes aegypti GN=AAEL005531 PE=4 SV=1
CCG006874.1	1619	436	139	11.5718371	3.001359155	-1.946930048	Down	1.51E-51	7.08E-51	AAEL013730-PA OS=Aedes aegypti GN=AAEL013730 PE=4 SV=1
CCG003708.1	2613	659	139	10.8369901	1.859625133	-2.542880378	Down	3.97E-107	2.13E-106	AAEL000334-PA OS=Aedes aegypti GN=AAEL000334 PE=4 SV=1
CCG014717.1	708	12	140	0.72830059	6.912653599	3.24663372	Up	1.30E-07	2.49E-07	-

CCG023502.1	1299	16	140	0.5292654	3.76763568	2.831596221	Up	7.09E-09	1.42E-08	AAEL010782-PA OS=Aedes aegypti GN=AAEL010782 PE=3 SV=1
CCG022896.1	982	24	140	1.05017681	4.983868379	2.24663372	Up	1.93E-11	4.23E-11	AAEL003567-PA (Fragment) OS=Aedes aegypti GN=AAEL003567 PE=4 SV=1
CCG001573.1	722	30	140	1.78544604	6.778613225	1.924705625	Up	4.55E-13	1.19E-12	AAEL001382-PA OS=Aedes aegypti GN=AAEL001382 PE=4 SV=1
CCG016651.1	825	37	140	1.92712749	5.932313634	1.622142855	Up	3.24E-11	7.02E-11	Mitochondrial 2-oxoglutarate/malate carrier protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016780 PE=3 SV=1
CCG010932.1	1318	56	140	1.82572469	3.713322267	1.024241299	Up	3.17E-06	5.70E-06	AAEL008250-PA OS=Aedes aegypti GN=AAEL008250 PE=4 SV=1
CCG010395.3	1731	255	140	6.33003023	2.827359184	-1.162757216	Down	3.76E-15	1.36E-14	AGAP007868-PA OS=Anopheles gambiae GN=AGAP007868 PE=4 SV=2
CCG014782.1	638	531	140	35.7632117	7.671095216	-2.22097183	Down	8.43E-74	4.22E-73	AAEL008307-PA OS=Aedes aegypti GN=AAEL008307 PE=4 SV=1
CCG015405.1	2157	563	140	11.2155589	2.268965576	-2.305394891	Down	8.69E-82	4.43E-81	AAEL005397-PA OS=Aedes aegypti GN=AAEL005397 PE=4 SV=1
CCG005452.1	972	719	140	31.7852255	5.035142745	-2.65825174	Down	1.04E-122	5.69E-122	AAEL001184-PA OS=Aedes aegypti GN=AAEL001184 PE=4 SV=1
CCG019323.1	1107	777	140	30.1603286	4.421100947	-2.770174568	Down	1.36E-138	7.62E-138	Prophenoloxidase activator OS=Aedes aegypti PE=2 SV=1
CCG010197.1	318	0	141	0.001	15.500368	13.92001485	Up	4.44E-16	1.63E-15	AAEL012697-PA OS=Aedes aegypti GN=AAEL012697 PE=4 SV=1
CCG023733.1	729	8	141	0.47154716	6.7614774	3.841864556	Up	2.19E-06	3.96E-06	AAEL014795-PA OS=Aedes aegypti GN=AAEL014795 PE=4 SV=1
CCG004962.1	1909	8	141	0.18007222	2.582041396	3.841864556	Up	2.19E-06	3.96E-06	AAEL010940-PA OS=Aedes aegypti GN=CYP314A1 PE=3 SV=2
CCG011369.1	1239	17	141	0.58957667	3.978302684	2.754401715	Up	3.39E-09	6.88E-09	AAEL005888-PA OS=Aedes aegypti GN=AAEL005888 PE=4 SV=1
CCG011801.1	691	21	141	1.30588195	7.133309732	2.449547133	Up	1.76E-10	3.73E-10	AAEL008656-PA OS=Aedes aegypti GN=AAEL008656 PE=4 SV=1

CCG010801.1	1449	51	141	1.51239232	3.401737077	1.169439214	Up	1.96E-07	3.72E-07	AAEL013796-PA OS=Aedes aegypti GN=AAEL013796 PE=4 SV=1
CCG027511.1	915	253	141	11.881249	5.387013142	-1.141129018	Down	1.22E-14	4.35E-14	AAEL005088-PA OS=Aedes aegypti GN=AAEL005088 PE=4 SV=1
CCG018666.1	2055	330	141	6.90024935	2.398597092	-1.524457658	Down	2.49E-28	1.04E-27	Ca-activated cation channel protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009092 PE=4 SV=1
CCG019418.1	252	341	141	58.1455536	19.55998819	-1.571763373	Down	1.70E-30	7.17E-30	AAEL012879-PA OS=Aedes aegypti GN=AAEL012879 PE=4 SV=1
CCG021560.1	1551	472	141	13.0765408	3.178025161	-2.040778493	Down	3.67E-59	1.76E-58	AAEL004951-PA OS=Aedes aegypti GN=AAEL004951 PE=4 SV=1
CCG004602.1	1779	2158	141	52.1240513	2.770723454	-4.233614593	Down	0	0	AAEL004341-PA (Fragment) OS=Aedes aegypti GN=AAEL004341 PE=4 SV=1
CCG013631.1	1731	6	142	0.14894189	2.86775003	4.267097823	Up	8.38E-06	1.47E-05	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC001426 PE=4 SV=1
CCG021642.1	638	16	142	1.0776109	7.780682291	2.852060323	Up	7.09E-09	1.42E-08	-
CCG013825.1	1494	21	142	0.60399225	3.322674231	2.459742901	Up	1.76E-10	3.72E-10	AAEL002819-PA OS=Aedes aegypti GN=AAEL002819 PE=4 SV=1
CCG007279.1	785	35	142	1.91584804	6.323662805	1.722777306	Up	3.17E-12	7.41E-12	-
CCG000448.1	2022	41	142	0.87129531	2.455032296	1.494508319	Up	3.01E-10	6.32E-10	AAEL002186-PA OS=Aedes aegypti GN=AAEL002186 PE=4 SV=1
CCG011755.1	876	41	142	2.01114055	5.666752627	1.494508319	Up	3.01E-10	6.32E-10	AAEL008243-PA OS=Aedes aegypti GN=AAEL008243 PE=4 SV=1
CCG025706.1	549	42	142	3.2873021	9.042031515	1.459742901	Up	6.02E-10	1.25E-09	AAEL013591-PA OS=Aedes aegypti GN=AAEL013591 PE=4 SV=1
CCG021442.1	168	44	142	11.2539781	29.54806727	1.392628705	Up	2.29E-09	4.67E-09	AAEL003534-PA OS=Aedes aegypti GN=AAEL003534 PE=4 SV=1
CCG002766.1	5112	50	142	0.42028301	0.971063244	1.208204134	Up	8.44E-08	1.62E-07	AAEL012505-PA OS=Aedes aegypti GN=AAEL012505 PE=4 SV=1
CCG012422.1	628	55	142	3.76327294	7.904578506	1.07070061	Up	1.14E-06	2.10E-06	-
CCG024327.1	549	278	142	21.7588091	9.042031515	-1.266880749	Down	1.49E-18	5.75E-18	AAEL008179-PA OS=Aedes aegypti GN=AAEL008179 PE=2 SV=1

CCG019230.1	1556	283	142	7.81518952	3.190279757	-1.29259792	Down	2.06E-19	7.99E-19	AAEL001757-PA OS=Aedes aegypti GN=AAEL001757 PE=4 SV=1
CCG019119.2	2814	292	142	4.45883529	1.764063718	-1.337764236	Down	5.43E-21	2.14E-20	AAEL002172-PA OS=Aedes aegypti GN=AAEL002172 PE=4 SV=1
CCG027056.1	3915	357	142	3.91831296	1.26796304	-1.627719941	Down	1.94E-33	8.34E-33	AAEL014591-PA OS=Aedes aegypti GN=AAEL014591 PE=4 SV=1
CCG020958.1	1074	373	142	14.9233808	4.622044043	-1.690971497	Down	9.68E-37	4.24E-36	AAEL008164-PA OS=Aedes aegypti GN=AAEL008164 PE=4 SV=1
CCG000543.1	1282	386	142	12.9378452	3.872133621	-1.740396714	Down	1.75E-39	7.77E-39	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005513 PE=4 SV=1
CCG027827.1	519	506	142	41.8934214	9.564692296	-2.130933251	Down	7.14E-67	3.51E-66	AAEL017144-PA OS=Aedes aegypti PE=4 SV=1
CCG015322.1	294	0	143	0.001	17.00351557	14.05354544	Up	4.44E-16	1.63E-15	AAEL013570-PA OS=Aedes aegypti GN=AAEL013570 PE=4 SV=1
CCG000908.1	513	9	143	0.75385499	9.744704831	3.692259539	Up	1.09E-06	2.01E-06	AAEL004861-PA OS=Aedes aegypti GN=AAEL004861 PE=4 SV=1
CCG001303.1	1482	14	143	0.40592192	3.373167057	3.054829619	Up	3.07E-08	6.01E-08	AAEL014400-PA OS=Aedes aegypti GN=AAEL014400 PE=4 SV=1
CCG009406.1	798	15	143	0.80770178	6.264453106	2.955293945	Up	1.48E-08	2.93E-08	AAEL002310-PA OS=Aedes aegypti GN=AAEL002310 PE=4 SV=1
CCG011366.1	3423	16	143	0.20085181	1.46042465	2.862184541	Up	7.09E-09	1.42E-08	AAEL005888-PA OS=Aedes aegypti GN=AAEL005888 PE=4 SV=1
CCG026988.1	1661	20	143	0.51739596	3.009652967	2.540256446	Up	3.69E-10	7.71E-10	AAEL010362-PA OS=Aedes aegypti GN=AAEL010362 PE=4 SV=1
CCG009133.1	531	31	143	2.50859091	9.414375854	1.90798823	Up	3.03E-13	8.37E-13	AAEL005626-PA OS=Aedes aegypti GN=AAEL005626 PE=4 SV=1
CCG011851.1	823	32	143	1.67075517	6.074159877	1.862184541	Up	3.35E-13	9.12E-13	-
CCG014676.1	662	48	143	3.1156303	7.551410239	1.27722204	Up	1.97E-08	3.89E-08	AAEL010795-PA OS=Aedes aegypti GN=AAEL010795 PE=4 SV=1
CCG010627.1	312	48	143	6.6107284	16.02254352	1.27722204	Up	1.97E-08	3.89E-08	AAEL002008-PA (Fragment) OS=Aedes aegypti GN=AAEL002008 PE=4 SV=1
CCG006803.1	1347	462	143	14.7379491	3.711235025	-1.989564501	Down	4.45E-56	2.12E-55	AAEL012248-PA OS=Aedes aegypti GN=AAEL012248 PE=3 SV=1

CCG024666.1	756	11	144	0.62522101	6.658719385	3.412806587	Up	2.67E-07	5.03E-07	Uncharacterized protein OS=Anopheles darlingi GN=AND_05262 PE=4 SV=1
CCG023148.1	1899	31	144	0.70145433	2.650864589	1.918041895	Up	2.82E-13	7.84E-13	AAEL003366-PA OS=Aedes aegypti GN=AAEL003366 PE=4 SV=1
CCG009036.1	1146	36	144	1.3498346	4.39266305	1.702313204	Up	3.37E-12	7.85E-12	-
CCG001219.2	1230	39	144	1.362455	4.092676305	1.586835986	Up	3.47E-11	7.52E-11	AAEL004974-PA OS=Aedes aegypti GN=AAEL004974 PE=4 SV=1
CCG001699.1	966	54	144	2.40203485	5.211171693	1.117350703	Up	3.96E-07	7.41E-07	AAEL003202-PA OS=Aedes aegypti GN=AAEL003202 PE=4 SV=1
CCG023239.1	756	56	144	3.1829433	6.658719385	1.064883283	Up	1.07E-06	1.97E-06	AAEL012822-PA OS=Aedes aegypti GN=AAEL012822 PE=4 SV=1
CCG012991.1	3387	289	144	3.66644621	1.486268632	-1.302687477	Down	5.16E-20	2.01E-19	AAEL004961-PA OS=Aedes aegypti GN=AAEL004961 PE=4 SV=1
CCG011789.1	1302	334	144	11.022958	3.866353191	-1.511466087	Down	2.60E-28	1.08E-27	AAEL013637-PA OS=Aedes aegypti GN=AAEL013637 PE=4 SV=1
CCG021319.1	1389	580	144	17.9427258	3.624184201	-2.307670885	Down	2.65E-84	1.35E-83	Ammonium transporter OS=Aedes albopictus PE=2 SV=1
CCG009039.1	381	1166	144	131.503177	13.21257705	-3.315113868	Down	3.04E-248	1.87E-247	AAEL002920-PA OS=Aedes aegypti GN=AAEL002920 PE=4 SV=1
CCG020603.1	2229	21	145	0.40482926	2.27409158	2.489904871	Up	1.76E-10	3.73E-10	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012788 PE=4 SV=1
CCG001296.1	903	28	145	1.33239487	5.613455296	2.074867372	Up	1.17E-12	2.85E-12	AAEL004611-PA OS=Aedes aegypti GN=AAEL004611 PE=4 SV=1
CCG005690.1	984	36	145	1.57206346	5.151372085	1.712297292	Up	2.33E-12	5.50E-12	AAEL009589-PA OS=Aedes aegypti GN=AAEL009589 PE=4 SV=1
CCG003142.1	470	50	145	4.57124836	10.78500028	1.238366104	Up	3.39E-08	6.64E-08	DNA-directed RNA polymerase subunit OS=Aedes aegypti GN=AAEL010468 PE=3 SV=1
CCG012560.1	1626	247	145	6.52738281	3.117435506	-1.066144938	Down	5.54E-13	1.42E-12	AAEL011699-PA OS=Aedes aegypti GN=AAEL011699 PE=4 SV=1
CCG026823.1	1254	279	145	9.56025196	4.042224986	-1.241899018	Down	4.38E-18	1.67E-17	AGAP007334-PA OS=Anopheles gambiae GN=AGAP007334 PE=4 SV=3

CCG013954.1	7323	409	145	2.399921	0.692195839	-1.793734739	Down	1.78E-43	8.05E-43	AAEL011854-PA OS=Aedes aegypti GN=AAEL011854 PE=4 SV=1
CCG006920.1	474	2	146	0.1813069	10.76773926	5.892137763	Up	9.36E-05	0.00015354	AAEL004436-PA OS=Aedes aegypti GN=AAEL004436 PE=4 SV=1
CCG017130.1	413	3	146	0.31212882	12.35813174	5.307175262	Up	5.28E-05	8.83E-05	-
CCG028032.1	1797	8	146	0.19129542	2.840238402	3.892137763	Up	2.19E-06	3.97E-06	AAEL000452-PA OS=Aedes aegypti GN=AAEL000452 PE=4 SV=1
CCG021284.1	1617	11	146	0.29231112	3.156405942	3.432706144	Up	2.67E-07	5.03E-07	AAEL011013-PA OS=Aedes aegypti GN=AAEL011013 PE=4 SV=1
CCG000777.1	1173	15	146	0.5494851	4.351158064	2.985247167	Up	1.48E-08	2.93E-08	AAEL000550-PA OS=Aedes aegypti GN=AAEL000550 PE=4 SV=1
CCG021747.1	1721	16	146	0.3994862	2.965664386	2.892137763	Up	7.09E-09	1.42E-08	Uncharacterized protein OS=Anopheles darlingi GN=AND_00142 PE=4 SV=1
CCG015718.1	263	24	146	3.92119251	19.40649585	2.307175262	Up	1.93E-11	4.23E-11	-
CCG019355.1	399	25	146	2.69233926	12.7917504	2.248281573	Up	9.32E-12	2.08E-11	AAEL013782-PA OS=Aedes aegypti GN=AAEL013782 PE=4 SV=1
CCG015177.1	1073	27	146	1.08125148	4.756671397	2.137250261	Up	2.28E-12	5.40E-12	Pol polyprotein OS=Drosophila melanogaster GN=pol PE=4 SV=1
CCG007329.1	2001	34	146	0.73012043	2.550678865	1.804674921	Up	3.73E-13	9.98E-13	AAEL000033-PA OS=Aedes aegypti GN=AAEL000033 PE=4 SV=1
CCG020718.1	2363	38	146	0.69100716	2.159927384	1.644210249	Up	7.78E-12	1.74E-11	AAEL003153-PA OS=Aedes aegypti GN=AAEL003153 PE=4 SV=1
CCG009890.1	1014	50	146	2.11882321	5.033440245	1.248281573	Up	2.50E-08	4.91E-08	AAEL005203-PA OS=Aedes aegypti GN=AAEL005203 PE=4 SV=1
CCG002907.1	986	51	146	2.22257248	5.176377697	1.219712421	Up	4.41E-08	8.58E-08	AAEL007974-PA OS=Aedes aegypti GN=AAEL007974 PE=4 SV=1
CCG026370.1	558	55	146	4.23536811	9.146789263	1.110778049	Up	3.73E-07	6.99E-07	AAEL014183-PA OS=Aedes aegypti GN=AAEL014183 PE=4 SV=1
CCG006624.1	1531	56	146	1.57172119	3.333708954	1.084782841	Up	6.15E-07	1.14E-06	Putative uncharacterized protein OS=Caenorhabditis brenneri GN=CAEBREN_31323 PE=4 SV=1
CCG008198.1	19131	266	146	0.59745698	0.26678733	-1.163144673	Down	9.53E-16	3.47E-15	Uncharacterized protein OS=Anopheles darlingi GN=AND_22381 PE=4 SV=1

CCG018574.1	1366	571	146	17.9617265	3.736389758	-2.265209173	Down	3.61E-81	1.84E-80	Out at first protein (Fragment) OS=Culex quinquefasciatus GN=CpipJ_CPIJ020077 PE=4 SV=1
CCG000290.1	606	0	147	0.001	8.479978029	13.04984481	Up	4.44E-16	1.65E-15	AAEL010818-PA OS=Aedes aegypti GN=AAEL010818 PE=3 SV=1
CCG015710.1	297	2	147	0.28935848	17.30258143	5.901985549	Up	9.36E-05	0.00015407	AAEL007013-PA OS=Aedes aegypti GN=AAEL007013 PE=4 SV=1
CCG025417.1	615	7	147	0.48908641	8.355880789	4.094630627	Up	4.31E-06	7.70E-06	-
CCG003369.1	1356	10	147	0.31688595	3.789724694	3.580057454	Up	5.43E-07	1.01E-06	AAEL001732-PA OS=Aedes aegypti GN=AAEL001732 PE=4 SV=1
CCG009001.1	1557	20	147	0.55195549	3.300492412	2.580057454	Up	3.69E-10	7.73E-10	Uncharacterized protein OS=Anopheles darlingi GN=AND_17136 PE=4 SV=1
CCG016264.1	301	22	147	3.14064505	17.0726468	2.44255393	Up	8.38E-11	1.80E-10	AAEL006177-PA OS=Aedes aegypti GN=AAEL006177 PE=4 SV=1
CCG026951.1	1989	50	147	1.08018438	2.583643381	1.258129359	Up	1.83E-08	3.62E-08	AAEL005053-PA OS=Aedes aegypti GN=AAEL005053 PE=4 SV=1
CCG001787.1	447	50	147	4.80645801	11.49634605	1.258129359	Up	1.83E-08	3.62E-08	AAEL009459-PA OS=Aedes aegypti GN=AAEL009459 PE=4 SV=1
CCG018511.1	1667	51	147	1.31461096	3.082703471	1.229560207	Up	3.25E-08	6.36E-08	-
CCG024619.1	2168	276	147	5.47031677	2.370325962	-1.206538908	Down	3.57E-17	1.35E-16	AAEL009038-PA (Fragment) OS=Aedes aegypti GN=AAEL009038 PE=4 SV=1
CCG023336.1	876	307	147	15.059028	5.866286171	-1.360109297	Down	1.61E-22	6.42E-22	AAEL002935-PA OS=Aedes aegypti GN=AAEL002935 PE=4 SV=1
CCG028363.1	1827	388	147	9.12548277	2.812734913	-1.697927294	Down	2.29E-38	1.02E-37	AAEL008940-PA OS=Aedes aegypti GN=AAEL008940 PE=4 SV=1
CCG008089.1	1876	412	147	9.43685003	2.739267956	-1.784514979	Down	1.76E-43	7.99E-43	AAEL006707-PA (Fragment) OS=Aedes aegypti GN=AAEL006707 PE=4 SV=1
CCG015347.1	2166	464	147	9.20496623	2.372514629	-1.955995446	Down	4.42E-55	2.10E-54	AAEL000474-PA OS=Aedes aegypti GN=AAEL000474 PE=4 SV=1
CCG012608.1	1911	5274	147	118.588373	2.689098213	-5.462696307	Down	0	0	AAEL009910-PA OS=Aedes aegypti GN=AAEL009910 PE=4 SV=1

CCG015185.1	1449	4	148	0.11861901	3.570617641	4.911766569	Up	2.95E-05	5.03E-05	AAEL014397-PA OS=Aedes aegypti GN=AAEL014397 PE=4 SV=1
CCG004487.1	834	17	148	0.87588188	6.203627053	2.824303728	Up	3.39E-09	6.89E-09	AAEL011589-PA OS=Aedes aegypti GN=AAEL011589 PE=4 SV=1
CCG007903.1	539	44	148	3.50773344	9.59893314	1.452334951	Up	3.04E-10	6.38E-10	Uncharacterized protein OS=Anopheles darlingi GN=AND_08991 PE=4 SV=1
CCG006114.1	878	52	148	2.54490456	5.892739137	1.211326851	Up	4.21E-08	8.21E-08	AAEL002303-PA OS=Aedes aegypti GN=AAEL002303 PE=4 SV=1
CCG020633.1	234	60	148	11.0178807	22.11036309	1.004875974	Up	2.39E-06	4.31E-06	Myosin heavy chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ015897 PE=4 SV=1
CCG015284.1	321	263	148	35.2057327	16.11783477	-1.12715242	Down	6.86E-15	2.47E-14	Actinin OS=Culex quinquefasciatus GN=CpipJ_CPIJ003874 PE=4 SV=1
CCG024878.1	2529	271	148	4.60450695	2.04579872	-1.170382472	Down	3.68E-16	1.38E-15	AAEL011941-PA OS=Aedes aegypti GN=AAEL011941 PE=4 SV=1
CCG021564.1	482	368	148	32.8067683	10.73407669	-1.611795387	Down	5.96E-34	2.57E-33	AAEL011100-PA OS=Aedes aegypti GN=AAEL011100 PE=4 SV=1
CCG000106.1	618	480	148	33.3745512	8.371885052	-1.995124026	Down	2.09E-58	1.00E-57	AAEL009032-PA OS=Aedes aegypti GN=AAEL009032 PE=4 SV=1
CCG015673.1	3660	527	148	6.18717217	1.413613378	-2.129892582	Down	1.56E-69	7.74E-69	AAEL001442-PA OS=Aedes aegypti GN=AAEL001442 PE=4 SV=1
CCG014082.1	1114	2	149	0.07714495	4.67574797	5.921481724	Up	9.36E-05	0.00015355	AAEL004775-PA OS=Aedes aegypti GN=AAEL004775 PE=4 SV=1
CCG023397.2	1128	13	149	0.49521857	4.617715637	3.221042006	Up	6.34E-08	1.23E-07	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015361 PE=4 SV=1
CCG012349.1	507	31	149	2.62734078	10.2737342	1.967285414	Up	2.45E-13	6.90E-13	AAEL000463-PA OS=Aedes aegypti GN=AAEL000463 PE=4 SV=1
CCG016727.1	1734	32	149	0.79298241	3.003911903	1.921481724	Up	1.91E-13	5.50E-13	AAEL001004-PA OS=Aedes aegypti GN=AAEL001004 PE=4 SV=1
CCG023585.1	3601	37	149	0.44151074	1.446482432	1.712028359	Up	1.18E-12	2.89E-12	-
CCG027014.1	2244	44	149	0.84254382	2.321204652	1.462050106	Up	2.16E-10	4.56E-10	AAEL007644-PA (Fragment) OS=Aedes aegypti GN=GPRNNA6 PE=3 SV=2

CCG005990.1	659	46	149	2.99940484	7.904071683	1.397919768	Up	8.22E-10	1.70E-09	AAEL007825-PA OS=Aedes aegypti GN=AAEL007825 PE=4 SV=1
CCG004916.1	1296	59	149	1.95618391	4.01912287	1.038838675	Up	1.15E-06	2.11E-06	AAEL012138-PA OS=Aedes aegypti GN=AAEL012138 PE=4 SV=1
CCG022320.1	972	246	149	10.8750563	5.358830493	-1.021032781	Down	3.79E-12	8.78E-12	AAEL003894-PA OS=Aedes aegypti GN=GNBPB5 PE=4 SV=1
CCG016210.1	393	268	149	29.3025162	13.25390137	-1.144607466	Down	1.73E-15	6.29E-15	AAEL002608-PA OS=Aedes aegypti GN=AAEL002608 PE=4 SV=1
CCG006497.1	1725	328	149	8.17047707	3.019584486	-1.43607028	Down	6.88E-26	2.82E-25	AAEL002367-PA (Fragment) OS=Aedes aegypti GN=AAEL002367 PE=4 SV=1
CCG007696.1	750	458	149	26.2401846	6.945044319	-1.917722064	Down	5.32E-53	2.51E-52	Steroid dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007736 PE=3 SV=1
CCG020243.1	1026	574	149	24.0395981	5.076786783	-2.243425202	Down	1.26E-80	6.42E-80	AAEL007060-PA OS=Aedes aegypti GN=AAEL007060 PE=3 SV=1
CCG005944.1	384	714	149	79.8968503	13.56453969	-2.55829854	Down	1.02E-116	5.53E-116	AAEL002860-PA OS=Aedes aegypti GN=AAEL002860 PE=4 SV=1
CCG003389.1	636	0	150	0.001	8.244876597	13.00928219	Up	4.44E-16	1.63E-15	AAEL010110-PA OS=Aedes aegypti GN=AAEL010110 PE=4 SV=1
CCG001048.1	1515	0	150	0.001	3.461215522	11.75706306	Up	4.44E-16	1.63E-15	Cytochrome P450 6CB1 (Fragment) OS=Aedes aegypti GN=CYP6CB1 PE=2 SV=1
CCG020723.1	2793	0	150	0.001	1.877458473	10.87456528	Up	4.44E-16	1.65E-15	AAEL011258-PA OS=Aedes aegypti GN=AAEL011258 PE=4 SV=1
CCG016470.1	1245	26	150	0.89735992	4.211840575	2.230692176	Up	4.60E-12	1.05E-11	-
CCG016241.3	1005	26	150	1.11165483	5.21765325	2.230692176	Up	4.60E-12	1.05E-11	-
CCG020705.1	756	30	150	1.7051482	6.936166026	2.024241299	Up	3.72E-13	9.96E-13	AAEL012822-PA OS=Aedes aegypti GN=AAEL012822 PE=4 SV=1
CCG018991.1	1178	34	150	1.24021305	4.451393477	1.843669053	Up	1.47E-13	4.39E-13	AAEL004775-PA OS=Aedes aegypti GN=AAEL004775 PE=4 SV=1
CCG010340.1	947	42	150	1.90573269	5.53721385	1.538814472	Up	3.72E-11	8.07E-11	AAEL003847-PA OS=Aedes aegypti GN=AAEL003847 PE=4 SV=1
CCG009770.1	498	54	150	4.65936881	10.52960144	1.176244392	Up	6.94E-08	1.34E-07	AAEL006748-PA OS=Aedes aegypti GN=AAEL006748 PE=4 SV=1

CCG019470.1	393	256	150	27.9904633	13.34285373	-1.068868106	Down	1.88E-13	5.44E-13	AAEL014640-PA OS=Aedes aegypti GN=PGRPLC PE=4 SV=1
CCG023032.1	591	290	150	21.0849798	8.872659079	-1.248777196	Down	6.89E-19	2.66E-18	AAEL003730-PA OS=Aedes aegypti GN=AAEL003730 PE=4 SV=1
CCG020054.1	1320	427	150	13.9000581	3.972531451	-1.806960365	Down	9.15E-46	4.19E-45	AAEL007796-PA OS=Aedes aegypti GN=CLIPD1 PE=3 SV=1
CCG004535.1	1533	619	150	17.3504669	3.420575027	-2.342663705	Down	1.69E-91	8.82E-91	Adenylylurate kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ006069 PE=3 SV=1
CCG013978.1	1170	7	151	0.25708388	4.511709224	4.133363021	Up	4.31E-06	7.69E-06	AAEL010401-PA OS=Aedes aegypti GN=AAEL010401 PE=4 SV=1
CCG017823.1	1674	8	151	0.20535118	3.153345157	3.940717943	Up	2.19E-06	3.97E-06	AAEL009254-PA OS=Aedes aegypti GN=AAEL009254 PE=3 SV=1
CCG003333.1	521	9	151	0.74227948	10.13186141	3.770792942	Up	1.09E-06	2.01E-06	-
CCG017780.1	885	10	151	0.48553372	5.964632534	3.618789848	Up	5.43E-07	1.01E-06	AAEL008915-PA OS=Aedes aegypti GN=AAEL008915 PE=4 SV=1
CCG012871.1	2863	35	151	0.52530238	1.843765209	1.811434926	Up	1.61E-13	4.76E-13	Retrovirus-related Pol polyprotein from transposon TNT 1-94 OS=Trichinella spiralis GN=Tsp_01017 PE=4 SV=1
CCG027717.1	1626	37	151	0.97778609	3.246432837	1.731264578	Up	5.82E-13	1.49E-12	AAEL006795-PA OS=Aedes aegypti GN=CYP9J15 PE=3 SV=2
CCG000360.1	960	41	151	1.83516575	5.498645617	1.583165939	Up	1.25E-11	2.78E-11	AAEL009174-PA OS=Aedes aegypti GN=AAEL009174 PE=2 SV=1
CCG025669.1	1245	48	151	1.65666447	4.239919512	1.355755442	Up	1.53E-09	3.15E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000155 PE=4 SV=1
CCG025732.1	1986	50	151	1.08181608	2.657955585	1.296861753	Up	5.25E-09	1.06E-08	AAEL004692-PA OS=Aedes aegypti GN=AAEL004692 PE=4 SV=1
CCG012009.1	3468	261	151	3.23388141	1.522116434	-1.087188053	Down	4.97E-14	1.68E-13	AAEL001304-PA OS=Aedes aegypti GN=AAEL001304 PE=3 SV=1
CCG007748.1	1221	285	151	10.0297906	4.323259453	-1.214100166	Down	7.61E-18	2.90E-17	AAEL000044-PA OS=Aedes aegypti GN=CYP325Q1 PE=3 SV=1
CCG011387.1	1824	346	151	8.15105711	2.894024009	-1.493910284	Down	8.61E-29	3.60E-28	AAEL014138-PA OS=Aedes aegypti GN=SRPN16 PE=3 SV=2

CCG008845.1	869	412	151	20.3723023	6.074453156	-1.745782584	Down	3.30E-42	1.49E-41	Fatty acid desaturase OS=Culex quinquefasciatus GN=CpipJ_CPIJ004097 PE=4 SV=1
CCG001744.1	1591	473	151	12.774786	3.317850278	-1.94497843	Down	1.06E-55	5.03E-55	AAEL004120-PA OS=Aedes aegypti GN=AAEL004120 PE=4 SV=1
CCG013624.1	468	561	151	51.5085921	11.27927306	-2.191139017	Down	1.42E-76	7.13E-76	AGAP012191-PA (Fragment) OS=Anopheles gambiae GN=AGAP012191 PE=4 SV=4
CCG009368.1	1914	772	151	17.3315753	2.757941375	-2.651739094	Down	2.92E-131	1.63E-130	AAEL014527-PA OS=Aedes aegypti GN=AAEL014527 PE=4 SV=1
CCG005067.1	384	814	151	91.0868853	13.74661404	-2.728167041	Down	1.03E-142	5.82E-142	AAEL002860-PA OS=Aedes aegypti GN=AAEL002860 PE=4 SV=1
CCG000022.1	1560	13	152	0.35808112	3.40619107	3.249800999	Up	6.34E-08	1.23E-07	AAEL003543-PA OS=Aedes aegypti GN=AAEL003543 PE=4 SV=1
CCG004905.1	432	20	152	1.98933957	12.30013442	2.628312622	Up	3.69E-10	7.71E-10	AAEL000408-PA OS=Aedes aegypti GN=AAEL000408 PE=4 SV=1
CCG003805.1	477	28	152	2.52233243	11.13974438	2.142885795	Up	1.17E-12	2.86E-12	AAEL013273-PA OS=Aedes aegypti GN=AAEL013273 PE=4 SV=1
CCG027934.1	1257	49	152	1.67503341	4.227253834	1.335530873	Up	2.07E-09	4.24E-09	AAEL005618-PA OS=Aedes aegypti GN=AAEL005618 PE=4 SV=1
CCG000304.1	1977	51	152	1.1084757	2.687738022	1.277815375	Up	6.95E-09	1.40E-08	AAEL000305-PA OS=Aedes aegypti GN=AAEL000305 PE=4 SV=1
CCG004927.1	3660	60	152	0.70442188	1.451819145	1.043350122	Up	8.20E-07	1.51E-06	AAEL013736-PA (Fragment) OS=Aedes aegypti GN=AAEL013736 PE=4 SV=1
CCG017513.1	1944	389	152	8.59836768	2.733363204	-1.653385628	Down	4.43E-37	1.94E-36	AAEL005160-PA OS=Aedes aegypti GN=AAEL005160 PE=4 SV=1
CCG013321.1	534	749	152	60.2702832	9.950670542	-2.598581191	Down	1.53E-124	8.43E-124	AAEL007659-PA OS=Aedes aegypti GN=AAEL007659 PE=4 SV=1
CCG008082.1	5973	771	152	5.54657046	0.889612936	-2.640346333	Down	1.83E-130	1.02E-129	AAEL000581-PA OS=Aedes aegypti GN=AAEL000581 PE=4 SV=1
CCG010696.1	363	808	152	95.646131	14.6381765	-2.707970765	Down	1.59E-140	8.95E-140	AAEL007184-PA OS=Aedes aegypti GN=AAEL007184 PE=4 SV=1
CCG011022.1	309	0	153	0.001	17.30943801	14.07927126	Up	4.44E-16	1.65E-15	AAEL013570-PA OS=Aedes aegypti GN=AAEL013570 PE=4 SV=1

CCG004373.1	885	2	153	0.09710674	6.043634289	5.959701047	Up	9.36E-05	0.00015367	-
CCG026909.1	1095	15	153	0.5886265	4.884581138	3.052810451	Up	1.48E-08	2.93E-08	AGAP004623-PA OS=Anopheles gambiae GN=AgaP_AGAP004623 PE=4 SV=5
CCG011590.1	315	17	153	2.31900155	16.97973443	2.872238205	Up	3.39E-09	6.88E-09	Uncharacterized protein OS=Anopheles darlingi GN=AND_13841 PE=4 SV=1
CCG024989.1	622	26	153	1.79616254	8.59906165	2.259261328	Up	4.60E-12	1.05E-11	-
CCG009441.1	1440	52	153	1.55168486	3.714316907	1.259261328	Up	9.15E-09	1.83E-08	AAEL005091-PA OS=Aedes aegypti GN=AAEL005091 PE=4 SV=1
CCG017464.1	1432	296	153	8.88201218	3.735067281	-1.249752319	Down	2.87E-19	1.11E-18	AAEL009480-PA OS=Aedes aegypti GN=AAEL009480 PE=4 SV=1
CCG024450.1	2643	908	153	14.7622092	2.023691391	-2.866847441	Down	1.10E-167	6.37E-167	AAEL005793-PA OS=Aedes aegypti GN=AAEL005793 PE=4 SV=1
CCG009385.1	1515	14	154	0.39708006	3.553514603	3.161744822	Up	3.07E-08	6.02E-08	AAEL003471-PA OS=Aedes aegypti GN=AAEL003471 PE=4 SV=1
CCG011067.1	525	14	154	1.14585959	10.25442785	3.161744822	Up	3.07E-08	6.00E-08	-
CCG010278.1	2087	31	154	0.6382663	2.579575766	2.014903434	Up	2.40E-13	6.77E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004626 PE=4 SV=1
CCG021604.1	2853	36	154	0.54220485	1.88698725	1.799174743	Up	1.33E-13	4.03E-13	AAEL005992-PA OS=Aedes aegypti GN=AAEL005992 PE=4 SV=1
CCG016162.1	946	47	154	2.13485996	5.690882265	1.414510893	Up	2.99E-10	6.28E-10	AAEL014058-PA OS=Aedes aegypti GN=AAEL014058 PE=4 SV=1
CCG004044.1	771	54	154	3.0095534	6.98258706	1.214212242	Up	2.10E-08	4.15E-08	AAEL009431-PA OS=Aedes aegypti GN=AAEL009431 PE=3 SV=1
CCG010174.2	1602	56	154	1.50206313	3.360533472	1.161744822	Up	6.25E-08	1.21E-07	AAEL006457-PA OS=Aedes aegypti GN=AAEL006457 PE=4 SV=1
CCG001133.1	1893	264	154	5.99260958	2.843937994	-1.075294375	Down	6.00E-14	1.99E-13	AAEL005762-PA OS=Aedes aegypti GN=AAEL005762 PE=4 SV=1
CCG015915.1	1022	316	154	13.286141	5.267685541	-1.334681004	Down	1.63E-22	6.51E-22	Voltage-dependent para-like sodium channel (Fragment) OS=Aedes albopictus PE=2 SV=1
CCG023119.1	243	376	154	66.488149	22.15462808	-1.585489107	Down	7.22E-34	3.11E-33	-

CCG020995.1	1374	648	154	20.2652023	3.918176581	-2.370750258	Down	4.46E-97	2.35E-96	AGAP008408-PA OS=Anopheles gambiae GN=AGAP008408 PE=4 SV=4
CCG020421.1	1146	1476	154	55.3432184	4.697709095	-3.558377262	Down	0	0	AAEL004058-PA OS=Aedes aegypti GN=AAEL004058 PE=4 SV=1
CCG000785.1	660	4	155	0.26042263	8.209898333	4.978437609	Up	2.95E-05	5.04E-05	AAEL007204-PA OS=Aedes aegypti GN=AAEL007204 PE=4 SV=1
CCG007918.1	921	16	155	0.74648833	5.883314766	2.978437609	Up	7.09E-09	1.42E-08	AAEL006871-PA OS=Aedes aegypti GN=AAEL006871 PE=4 SV=1
CCG008704.1	1629	19	155	0.50118168	3.326293984	2.730510096	Up	7.73E-10	1.60E-09	Synaptic vesicular amine transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ012480 PE=4 SV=1
CCG023123.1	690	27	155	1.6814244	7.852946231	2.223550107	Up	2.28E-12	5.41E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008409 PE=4 SV=1
CCG026328.1	2092	28	155	0.57512073	2.590120889	2.171082687	Up	1.17E-12	2.85E-12	AAEL010940-PA OS=Aedes aegypti GN=CYP314A1 PE=3 SV=2
CCG002099.1	333	32	155	4.12922375	16.27187057	1.978437609	Up	1.78E-13	5.19E-13	AAEL006133-PA OS=Aedes aegypti GN=AAEL006133 PE=4 SV=1
CCG012245.1	632	38	155	2.58362328	8.573628006	1.730510096	Up	3.08E-13	8.48E-13	-
CCG023086.1	1335	42	155	1.35185682	4.058826142	1.586120186	Up	6.30E-12	1.42E-11	AAEL010476-PA (Fragment) OS=Aedes aegypti GN=AAEL010476 PE=4 SV=1
CCG021479.1	2571	43	155	0.71866923	2.107558499	1.552172854	Up	1.32E-11	2.91E-11	AGAP005207-PA OS=Anopheles gambiae GN=AGAP005207 PE=4 SV=3
CCG019366.1	1677	253	155	6.48261351	3.231087	-1.004555966	Down	3.69E-12	8.56E-12	AAEL005178-PA (Fragment) OS=Aedes aegypti GN=AAEL005178 PE=4 SV=1
CCG018667.1	1488	258	155	7.45039753	3.641487164	-1.032789646	Down	6.97E-13	1.76E-12	Ca-activated Cl channel protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009092 PE=4 SV=1

CCG022909.1	1125	267	155	10.1981503	4.816473689	-1.082258323	Down	3.18E-14	1.10E-13	Rho/rac/cdc GTPase-activating protein OS=Culex quinquefasciatus AAEL008471-PA OS=Aedes aegypti GN=AAEL008471 PE=4 SV=1
CCG027762.1	4128	326	155	3.39344319	1.312629094	-1.370290545	Down	4.52E-24	1.82E-23	AAEL011752-PA OS=Aedes aegypti GN=GST11 PE=3 SV=2
CCG015905.1	1213	974	155	34.5033153	4.46705103	-2.949340353	Down	3.84E-185	2.26E-184	-
CCG025978.1	264	2	156	0.32552829	20.65716355	5.987715423	Up	9.36E-05	0.00015405	AAEL008458-PA OS=Aedes aegypti GN=AAEL008458 PE=4 SV=1
CCG007355.1	186	3	156	0.69306024	29.31984503	5.402752922	Up	5.28E-05	8.84E-05	CEC-A OS=Anopheles melas GN=CEC-A PE=4 SV=1
CCG022261.1	253	4	156	0.67936339	21.55530109	4.987715423	Up	2.95E-05	5.03E-05	AAEL009254-PA OS=Aedes aegypti GN=AAEL009254 PE=3 SV=1
CCG015236.1	867	18	156	0.89210522	6.290070561	2.817790421	Up	1.62E-09	3.33E-09	Gastfura zinc finger protein XLCGF62.1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007367 PE=4 SV=1
CCG002267.1	1008	24	156	1.02308892	5.4102095	2.402752922	Up	1.93E-11	4.23E-11	AAEL010250-PA OS=Aedes aegypti GN=AAEL010250 PE=4 SV=1
CCG023115.1	719	34	156	2.03194851	7.584827784	1.900252581	Up	9.33E-14	2.94E-13	AAEL005976-PA OS=Aedes aegypti GN=AAEL005976 PE=2 SV=1
CCG009801.1	552	37	156	2.88021772	9.879513001	1.778262057	Up	1.29E-13	3.96E-13	AAEL012227-PA OS=Aedes aegypti GN=AAEL012227 PE=4 SV=1
CCG004889.1	729	40	156	2.35773578	7.480783507	1.665787328	Up	9.90E-13	2.45E-12	AAEL006704-PA OS=Aedes aegypti GN=AAEL006704 PE=4 SV=1
CCG018059.1	1059	41	156	1.66360634	5.149661168	1.630163418	Up	2.12E-12	5.05E-12	AAEL001996-PA OS=Aedes aegypti GN=AAEL001996 PE=4 SV=1
CCG014718.1	2015	44	156	0.93829693	2.706447234	1.528283804	Up	1.90E-11	4.18E-11	AAEL010569-PA OS=Aedes aegypti GN=AAEL010569 PE=4 SV=1
CCG014304.1	1113	46	156	1.77592794	4.899812378	1.464153467	Up	7.75E-11	1.66E-10	GG13338 OS=Drosophila erecta GN=GG13338 PE=4 SV=1
CCG023036.1	1891	48	156	1.09071775	2.883919184	1.402752922	Up	2.96E-10	6.22E-10	AAEL007321-PA OS=Aedes aegypti GN=AAEL007321 PE=4 SV=1
CCG020388.1	1488	52	156	1.50163051	3.664980629	1.287275705	Up	3.59E-09	7.27E-09	

CCG005925.1	1536	391	156	10.9382593	3.550449985	-1.623309375	Down	2.51E-36	1.10E-35	AAEL004048-PA OS=Aedes aegypti GN=AAEL004048 PE=4 SV=1
CCG016460.1	873	657	156	32.3380477	6.246839836	-2.372034138	Down	1.83E-98	9.67E-98	AAEL004251-PA OS=Aedes aegypti GN=AAEL004251 PE=4 SV=1
CCG019020.1	642	5	157	0.33465525	8.548986687	4.675005858	Up	1.59E-05	2.76E-05	AAEL008029-PA OS=Aedes aegypti GN=AAEL008029 PE=3 SV=1
CCG022425.1	852	6	157	0.30260376	6.441842081	4.411971452	Up	8.38E-06	1.48E-05	Techniquein-55 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014772 PE=4 SV=1
CCG006533.1	1617	8	157	0.21258991	3.394217349	3.996933953	Up	2.19E-06	3.97E-06	AAEL001053-PA OS=Aedes aegypti GN=AAEL001053 PE=4 SV=1
CCG018201.1	332	8	157	1.03541529	16.53147426	3.996933953	Up	2.19E-06	3.97E-06	Cell division protein kinase 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001155 PE=4 SV=1
CCG018601.1	714	15	157	0.90272552	7.686903996	3.090043357	Up	1.48E-08	2.93E-08	AAEL006106-PA OS=Aedes aegypti GN=AAEL006106 PE=4 SV=1
CCG004322.1	5726	20	157	0.15008639	0.958513701	2.675005858	Up	3.69E-10	7.73E-10	-
CCG021151.1	2820	40	157	0.60949978	1.946258671	1.675005858	Up	7.04E-13	1.77E-12	AAEL011173-PA OS=Aedes aegypti GN=AAEL011173 PE=4 SV=1
CCG000918.1	1069	44	157	1.76863267	5.134190321	1.537502334	Up	1.34E-11	2.96E-11	AAEL007497-PA OS=Aedes aegypti GN=AAEL007497 PE=4 SV=1
CCG014648.1	1027	44	157	1.84096234	5.344157209	1.537502334	Up	1.34E-11	2.96E-11	AAEL004220-PA OS=Aedes aegypti GN=AAEL004220 PE=4 SV=1
CCG002453.1	1200	46	157	1.64717316	4.573707878	1.473371997	Up	5.50E-11	1.18E-10	AAEL011248-PA OS=Aedes aegypti GN=AAEL011248 PE=4 SV=1
CCG008531.1	1065	54	157	2.17874711	5.153473665	1.242046451	Up	8.45E-09	1.69E-08	AAEL002469-PA OS=Aedes aegypti GN=AAEL002469 PE=4 SV=1
CCG018471.1	1026	463	157	19.3908257	5.349365939	-1.857934431	Down	2.33E-51	1.09E-50	AAEL007164-PA OS=Aedes aegypti GN=AAEL007164 PE=4 SV=1
CCG021854.1	1062	718	157	29.0511012	5.1680315	-2.490906081	Down	8.24E-114	4.45E-113	AAEL005979-PA (Fragment) OS=Aedes aegypti GN=SCR3 PE=4 SV=2
CCG002765.1	1698	862	157	21.8138464	3.232302387	-2.754610106	Down	1.29E-152	7.35E-152	AAEL012509-PA (Fragment) OS=Aedes aegypti GN=AAEL012509 PE=4 SV=1

CCG015786.1	1134	6	158	0.22735309	4.870729921	4.421131451	Up	8.38E-06	1.47E-05	AAEL007080-PA OS=Aedes aegypti GN=AAEL007686 PE=4 SV=1
CCG022535.1	954	7	158	0.31529155	5.789735566	4.19873903	Up	4.31E-06	7.70E-06	AAEL000643-PA OS=Aedes aegypti GN=AAEL006643 PE=4 SV=1
CCG017888.1	480	14	158	1.25328393	11.50709944	3.19873903	Up	3.07E-08	6.01E-08	AAEL013597-PA OS=Aedes aegypti GN=AAEL013597 PE=4 SV=1
CCG024173.1	1449	32	158	0.94895204	3.81187559	2.006093952	Up	1.77E-13	5.16E-13	AAEL015402-PA OS=Aedes aegypti GN=AAEL015402 PE=4 SV=1
CCG026425.1	1761	37	158	0.90282804	3.136517734	1.796640586	Up	8.70E-14	2.77E-13	AAEL001146-PA OS=Aedes aegypti GN=AAEL001146 PE=3 SV=1
CCG015469.1	2115	38	158	0.77203306	2.611540298	1.758166439	Up	1.27E-13	3.90E-13	N-acetylglucosaminyltransferase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005697 PE=3 SV=1
CCG015844.1	5040	45	158	0.38365834	1.095914232	1.514240856	Up	1.92E-11	4.23E-11	AAEL003012-PA OS=Aedes aegypti GN=AAEL003012 PE=4 SV=1
CCG005359.1	903	52	158	2.47444762	6.116730598	1.305654234	Up	1.91E-09	3.91E-09	AAEL010800-PA OS=Aedes aegypti GN=AAEL010800 PE=4 SV=1
CCG023172.1	1581	316	158	8.58851116	3.493616528	-1.297686796	Down	1.29E-21	5.09E-21	AAEL007080-PA OS=Aedes aegypti GN=AAEL007088 PE=4 SV=1
CCG004203.1	1695	415	158	10.5206135	3.258647628	-1.690873574	Down	1.10E-40	4.92E-40	AAEL000780-PA OS=Aedes aegypti GN=AAEL000780 PE=4 SV=1
CCG025773.1	3891	468	158	5.16829499	1.419534241	-1.864270768	Down	3.94E-52	1.85E-51	AAEL005859-PA OS=Aedes aegypti GN=AAEL005859 PE=4 SV=1
CCG000091.1	3789	570	158	6.46417227	1.457748147	-2.148724157	Down	6.75E-76	3.39E-75	AGAP013009-PA OS=Anopheles gambiae GN=AgaP_AGAP013009 PE=4 SV=1
CCG014324.1	804	0	159	0.001	6.913390556	12.75517771	Up	4.44E-16	1.66E-15	AAEL009778-PA OS=Aedes aegypti GN=AAEL009778 PE=4 SV=1
CCG024054.1	1143	7	159	0.26315673	4.862962386	4.207841237	Up	4.31E-06	7.70E-06	AAEL013016-PA OS=Aedes aegypti GN=AAEL013016 PE=4 SV=1
CCG009508.1	2028	13	159	0.27544702	2.74081164	3.314756441	Up	6.34E-08	1.23E-07	AAEL001370-PA OS=Aedes aegypti GN=AAEL001370 PE=3 SV=1
CCG014434.1	612	23	159	1.61487565	9.082297397	2.491634203	Up	4.01E-11	8.67E-11	AAEL001738-PA OS=Aedes aegypti GN=AAEL001738 PE=4 SV=1

CCG004819.1	369	23	159	2.67833034	15.06332251	2.491634203	Up	4.01E-11	8.68E-11	AAEL008426-PA OS=Aedes aegypti GN=AAEL008426 PE=4 SV=1
CCG012047.1	884	46	159	2.23598167	6.287744352	1.491634203	Up	2.76E-11	6.01E-11	AAEL003050-PA OS=Aedes aegypti GN=AAEL003050 PE=4 SV=1
CCG002426.1	502	292	159	24.9943476	11.07244224	-1.1746284	Down	2.19E-17	8.31E-17	Uncharacterized protein OS=Anopheles darlingi GN=AND_07716 PE=4 SV=1
CCG028089.1	1716	309	159	7.73755711	3.23914103	-1.256266869	Down	3.37E-20	1.32E-19	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008574 PE=4 SV=1
CCG001708.1	963	335	159	14.9479347	5.771927318	-1.372821126	Down	9.30E-25	3.77E-24	AAEL010049-PA OS=Aedes aegypti GN=AAEL010049 PE=4 SV=1
CCG015496.1	1164	506	159	18.6792833	4.775228528	-1.967797416	Down	2.21E-60	1.07E-59	AAEL005781-PA OS=Aedes aegypti GN=AAEL005781 PE=4 SV=1
CCG008872.1	1166	648	159	23.8802642	4.767037742	-2.324653844	Down	8.18E-95	4.29E-94	AAEL009217-PA OS=Aedes aegypti GN=AAEL009217 PE=4 SV=1
CCG026979.1	1818	659	159	15.5759379	3.057407044	-2.348938496	Down	1.32E-97	6.94E-97	AAEL008625-PA OS=Aedes aegypti GN=AAEL008625 PE=3 SV=1
CCG019802.1	745	984	159	56.7546562	7.460893969	-2.927318346	Down	1.41E-185	8.32E-185	AAEL007762-PA OS=Aedes aegypti GN=AAEL007762 PE=4 SV=1
CCG002965.1	390	0	160	0.001	14.34185714	13.80794423	Up	4.44E-16	1.66E-15	AAEL011472-PA OS=Aedes aegypti GN=AAEL011472 PE=4 SV=1
CCG022834.1	1617	7	160	0.18601617	3.459075005	4.216886377	Up	4.31E-06	7.70E-06	AAEL006712-PA OS=Aedes aegypti GN=AAEL006712 PE=4 SV=1
CCG006182.1	600	15	160	1.07424337	9.322207139	3.117350703	Up	1.48E-08	2.92E-08	-
CCG012160.1	1002	29	160	1.24363503	5.582159964	2.166260304	Up	6.66E-13	1.69E-12	AAEL002712-PA OS=Aedes aegypti GN=AAEL002712 PE=4 SV=1
CCG005372.1	906	31	160	1.47026686	6.173647112	2.070044988	Up	2.40E-13	6.77E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000877 PE=4 SV=1
CCG011518.1	1425	40	160	1.20616799	3.925139848	1.702313204	Up	2.75E-13	7.66E-13	AAEL011900-PA OS=Aedes aegypti GN=AAEL011900 PE=4 SV=1
CCG019842.1	1191	46	160	1.65962031	4.696326015	1.500679343	Up	1.95E-11	4.28E-11	AAEL005363-PA (Fragment) OS=Aedes aegypti GN=AAEL005363 PE=4 SV=1

CCG020596.1	1203	54	160	1.92881602	4.64947987	1.269353797	Up	3.35E-09	6.81E-09	Krüppel protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014712 PE=4 SV=1
CCG026505.1	1380	338	160	10.5244712	4.053133539	-1.376638138	Down	4.59E-25	1.86E-24	AAEL002397-PA OS=Aedes aegypti GN=AAEL002397 PE=4 SV=1
CCG024547.1	1023	449	160	18.8596391	5.46757017	-1.786330336	Down	2.69E-47	1.24E-46	AGAP005912-PA OS=Anopheles gambiae GN=AGAP005912 PE=4 SV=3
CCG005886.1	1350	518	160	16.4876463	4.143203173	-1.992566989	Down	8.71E-63	4.23E-62	AAEL012101-PA OS=Aedes aegypti GN=AAEL012101 PE=3 SV=1
CCG017033.1	3378	2023	160	25.733503	1.655809439	-3.958039306	Down	0	0	Insulin receptor tyrosine kinase substrate OS=Culex quinquefasciatus GN=CpipJ_CPIJ007108 PE=4 SV=1
CCG013395.1	1755	2727	160	66.7683569	3.187079364	-4.388857686	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_09568 PE=4 SV=1
CCG011364.1	645	8	161	0.53295795	8.726019473	4.033230082	Up	2.19E-06	3.96E-06	AAEL005890-PA OS=Aedes aegypti GN=AAEL005890 PE=4 SV=1
CCG006241.1	627	21	161	1.43917771	8.976527209	2.640912659	Up	1.76E-10	3.73E-10	RPA-interacting protein alpha OS=Culex quinquefasciatus GN=CpipJ_CPIJ008848 PE=4 SV=1
CCG001224.3	1728	25	161	0.62166861	3.257107963	2.389373892	Up	9.32E-12	2.08E-11	AAEL003549-PA OS=Aedes aegypti GN=AAEL003549 PE=4 SV=1
CCG002836.1	672	46	161	2.94138064	8.375420477	1.509668126	Up	1.38E-11	3.05E-11	AAEL000279-PA (Fragment) OS=Aedes aegypti GN=AAEL000279 PE=3 SV=1
CCG012779.1	900	59	161	2.81690482	6.253647289	1.150587033	Up	3.98E-08	7.76E-08	AAEL005665-PA (Fragment) OS=Aedes aegypti GN=SRPN3 PE=3 SV=2
CCG009665.1	2052	283	161	5.92613786	2.742827758	-1.111428161	Down	1.39E-15	5.04E-15	AAEL010952-PA OS=Aedes aegypti GN=AAEL010952 PE=4 SV=1
CCG024737.1	2514	320	161	5.46949685	2.238775879	-1.288698013	Down	1.19E-21	4.71E-21	AAEL003741-PA OS=Aedes aegypti GN=AAEL003741 PE=4 SV=1

CCG000723.1	1512	366	161	10.401404	3.722409101	-1.482469756	Down	4.95E-30	2.08E-29	Voltage-dependent para-like sodium channel (Fragment) OS=Aedes albopictus PE=2 SV=1
CCG000402.1	960	719	161	32.1825408	5.862794334	-2.456617879	Down	3.87E-112	2.09E-111	AAEL007213-PA OS=Aedes aegypti GN=AAEL007213 PE=3 SV=1
CCG024223.1	1285	1289	161	43.1034925	4.379986428	-3.298806466	Down	5.77E-273	3.59E-272	-
CCG007962.1	2772	2	162	0.03100269	2.043016175	6.042163207	Up	9.36E-05	0.00015391	AAEL009308-PA OS=Aedes aegypti GN=AAEL009308 PE=4 SV=1
CCG004424.1	948	2	162	0.09065345	5.97388274	6.042163207	Up	9.36E-05	0.00015392	Putative salivary protein 34k-1 OS=Aedes albopictus PE=2 SV=1
CCG027104.1	330	9	162	1.17190185	17.16133587	3.872238205	Up	1.09E-06	2.01E-06	AAEL014451-PA OS=Aedes aegypti GN=AAEL014451 PE=4 SV=1
CCG000588.1	544	18	162	1.42179269	10.41036919	2.872238205	Up	1.62E-09	3.32E-09	Histone transcription regulator OS=Culex quinquefasciatus GN=CpipJ_CPIJ002429 PE=4 SV=1
CCG022825.1	567	20	162	1.51568729	9.988079078	2.720235112	Up	3.69E-10	7.71E-10	AAEL012187-PA OS=Aedes aegypti GN=AAEL012187 PE=4 SV=1
CCG023931.1	1668	44	162	1.1334942	3.39522832	1.582731588	Up	2.28E-12	5.41E-12	AAEL001475-PA OS=Aedes aegypti GN=AAEL001475 PE=4 SV=1
CCG015399.1	351	52	162	6.36588661	16.13458928	1.341723489	Up	5.31E-10	1.11E-09	AAEL004276-PA OS=Aedes aegypti GN=AAEL004276 PE=4 SV=1
CCG027758.1	1566	289	162	7.92991909	3.616373459	-1.132762476	Down	2.48E-16	9.30E-16	AAEL000714-PA OS=Aedes aegypti GN=AAEL000714 PE=4 SV=1
CCG026547.1	283	289	162	43.8807537	20.01145172	-1.132762476	Down	2.48E-16	9.30E-16	Aquaporin-1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ015704 PE=4 SV=1
CCG023276.2	378	342	162	38.8773789	14.98211862	-1.375689308	Down	2.56E-25	1.05E-24	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009959 PE=4 SV=1
CCG013181.1	1110	389	162	15.0587629	5.102018772	-1.561463138	Down	3.06E-34	1.32E-33	AAEL003614-PA OS=Aedes aegypti GN=CLIPB40 PE=3 SV=1
CCG025398.1	1785	514	162	12.3733578	3.172683942	-1.963461342	Down	3.93E-61	1.90E-60	AAEL005829-PA OS=Aedes aegypti GN=AAEL005829 PE=4 SV=1
CCG025113.1	558	17	163	1.30911378	10.21182637	2.963578517	Up	3.39E-09	6.88E-09	-

CCG006907.1	1032	60	163	2.49824038	5.521510769	1.144150762	Up	3.76E-08	7.35E-08	AAEL003956-PA OS=Aedes aegypti GN=AAEL003956 PE=4 SV=1
CCG002034.1	2214	323	163	6.26884565	2.573712337	-1.284348997	Down	9.87E-22	3.91E-21	AAEL001299-PA OS=Aedes aegypti GN=AAEL001299 PE=4 SV=1
CCG003407.1	1314	486	163	15.8929155	4.336529006	-1.873771146	Down	1.95E-54	9.23E-54	AAEL003781-PA OS=Aedes aegypti GN=AAEL003781 PE=3 SV=1
CCG009334.1	2823	628	163	9.55897745	2.018490653	-2.243579391	Down	4.76E-88	2.46E-87	AAEL007535-PA OS=Aedes aegypti GN=AAEL007535 PE=4 SV=1
CCG002637.1	1323	977	163	31.731996	4.307028809	-2.881173394	Down	3.11E-181	1.82E-180	AAEL014436-PA (Fragment) OS=Aedes aegypti GN=AAEL014436 PE=4 SV=1
CCG014123.1	645	0	164	0.001	8.88861611	13.1177431	Up	4.44E-16	1.63E-15	AAEL004817-PA OS=Aedes aegypti GN=AAEL004817 PE=4 SV=1
CCG028111.1	1725	0	164	0.001	3.323569502	11.69851781	Up	4.44E-16	1.64E-15	Putative 62 kDa salivary secreted protein OS=Aedes albopictus PE=2 SV=1
CCG026730.1	285	21	164	3.16619097	20.11634172	2.667547786	Up	1.76E-10	3.72E-10	AAEL003821-PA OS=Aedes aegypti GN=AAEL003821 PE=4 SV=1
CCG012535.2	480	27	164	2.41704757	11.9440779	2.304977706	Up	2.28E-12	5.40E-12	AAEL012816-PA OS=Aedes aegypti GN=AAEL012816 PE=4 SV=1
CCG014450.1	1005	28	164	1.19716674	5.70463422	2.252510286	Up	1.17E-12	2.85E-12	AAEL013879-PA OS=Aedes aegypti GN=AAEL013879 PE=4 SV=1
CCG024076.1	450	33	164	3.15111387	12.74034976	2.015471089	Up	1.31E-13	4.00E-13	AAEL006555-PA OS=Aedes aegypti GN=AAEL006555 PE=4 SV=1
CCG027510.1	965	37	164	1.64754423	5.941095742	1.850411843	Up	5.48E-14	1.83E-13	AAEL005229-PA OS=Aedes aegypti GN=AAEL005229 PE=4 SV=1
CCG002475.1	474	42	164	3.80744484	12.09526876	1.667547786	Up	3.07E-13	8.46E-13	AAEL009769-PA OS=Aedes aegypti GN=AAEL009769 PE=4 SV=1
CCG007364.1	1817	43	164	1.01689521	3.155287502	1.633600454	Up	5.91E-13	1.52E-12	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1
CCG018669.1	1836	45	164	1.05317977	3.122634744	1.568012112	Up	2.27E-12	5.40E-12	AAEL009724-PA OS=Aedes aegypti GN=AAEL009724 PE=4 SV=1
CCG008437.1	1314	297	164	9.71233728	4.363133478	-1.154453912	Down	3.25E-17	1.23E-16	AAEL010590-PA OS=Aedes aegypti GN=AAEL010590 PE=4 SV=1

CCG025599.1	1049	302	164	12.3706958	5.465354996	-1.178539531	Down	5.12E-18	1.96E-17	AAEL002512-PA OS=Aedes aegypti GN=AAEL002512 PE=4 SV=1
CCG014617.1	2112	313	164	6.36814722	2.714563158	-1.230153638	Down	7.99E-20	3.11E-19	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005107 PE=4 SV=1
CCG016988.1	332	764	164	98.8821604	17.26854636	-2.51756362	Down	1.74E-122	9.58E-122	-
CCG017554.1	933	814	164	37.4891361	6.144863227	-2.609019776	Down	8.39E-136	4.69E-135	AAEL008912-PA OS=Aedes aegypti GN=AAEL008912 PE=4 SV=1
CCG025842.2	1506	19	165	0.54211485	3.830090085	2.820707905	Up	7.73E-10	1.60E-09	AAEL008914-PA OS=Aedes aegypti GN=AAEL008914 PE=4 SV=1
CCG006455.1	747	20	165	1.15046144	7.721707721	2.746707323	Up	3.69E-10	7.72E-10	AGAP011122-PA OS=Anopheles gambiae GN=AGAP011122 PE=3 SV=4
CCG009384.1	987	21	165	0.91424967	5.844088822	2.676317995	Up	1.76E-10	3.73E-10	AAEL003484-PA OS=Aedes aegypti GN=AAEL003484 PE=4 SV=1
CCG007809.1	1353	22	165	0.69869487	4.263204484	2.609203799	Up	8.38E-11	1.80E-10	AAEL012306-PA OS=Aedes aegypti GN=AAEL012306 PE=4 SV=1
CCG023303.1	666	23	165	1.48393978	8.660834335	2.545073462	Up	4.01E-11	8.66E-11	AAEL001131-PA OS=Aedes aegypti GN=AAEL001131 PE=4 SV=1
CCG002119.1	1551	25	165	0.69261339	3.718965614	2.424779228	Up	9.32E-12	2.08E-11	AAEL003879-PA OS=Aedes aegypti GN=AAEL003879 PE=4 SV=1
CCG006639.1	1422	55	165	1.66197989	4.056340132	1.287275705	Up	1.27E-09	2.61E-09	AAEL014413-PA OS=Aedes aegypti GN=CYP304C1 PE=3 SV=1
CCG005339.1	1138	58	165	2.19002162	5.068642942	1.210654423	Up	7.10E-09	1.42E-08	AAEL006285-PA OS=Aedes aegypti GN=AAEL006285 PE=4 SV=1
CCG016739.1	870	295	165	14.5701974	6.630018009	-1.135935726	Down	1.04E-16	3.93E-16	AAEL000953-PA OS=Aedes aegypti GN=AAEL000953 PE=4 SV=1
CCG020845.1	450	337	165	32.1795568	12.81803482	-1.327969363	Down	9.92E-24	3.99E-23	26S proteasome regulatory complex subunit RPN5/PSMD12 OS=Aedes aegypti GN=AAEL014325 PE=2 SV=1
CCG024485.1	1905	668	165	15.0676025	3.02788224	-2.315068874	Down	3.39E-97	1.78E-96	AAEL010268-PA OS=Aedes aegypti GN=AAEL010268 PE=4 SV=1
CCG002812.1	964	7	166	0.31202089	6.019786249	4.269997713	Up	4.31E-06	7.69E-06	AAEL006009-PA OS=Aedes aegypti GN=AAEL006009 PE=4 SV=1

CCG022235.1	651	14	166	0.92408031	8.91409208	3.269997713	Up	3.07E-08	6.01E-08	AAEL013578-PA OS=Aedes aegypti GN=AAEL013578 PE=4 SV=1
CCG014734.1	892	30	166	1.44517045	6.505688278	2.17046204	Up	3.71E-13	9.95E-13	AAEL006838-PA OS=Aedes aegypti GN=AAEL006838 PE=4 SV=1
CCG019143.1	2116	56	166	1.13719524	2.742473509	1.269997713	Up	1.67E-09	3.42E-09	AAEL006381-PA OS=Aedes aegypti GN=AAEL006381 PE=4 SV=1
CCG006651.1	1008	59	166	2.51509359	5.757017802	1.194709586	Up	9.14E-09	1.83E-08	AAEL005746-PA OS=Aedes aegypti GN=AAEL005746 PE=4 SV=1
CCG012880.1	915	60	166	2.81768752	6.342157316	1.17046204	Up	1.57E-08	3.11E-08	AAEL011624-PA OS=Aedes aegypti GN=AAEL011624 PE=3 SV=1
CCG013350.1	976	62	166	2.72963478	5.945772484	1.123156325	Up	4.48E-08	8.71E-08	AAEL008250-PA OS=Aedes aegypti GN=AAEL008250 PE=4 SV=1
CCG012892.1	1184	67	166	2.43156437	4.901244885	1.011263445	Up	5.04E-07	9.40E-07	AAEL009499-PA OS=Aedes aegypti GN=AAEL009499 PE=4 SV=1
CCG019885.1	1461	451	166	13.2644424	3.971987641	-1.739630988	Down	8.05E-46	3.69E-45	AAEL009920-PA OS=Aedes aegypti GN=AAEL009920 PE=4 SV=1
CCG009309.1	828	742	166	38.5066945	7.008543411	-2.457922741	Down	9.71E-116	5.27E-115	Putative salivary secreted protein 275 OS=Aedes albopictus PE=2 SV=1
CCG026878.1	1062	787	166	31.8429201	5.464288083	-2.54286719	Down	1.41E-127	7.81E-127	AAEL010068-PA OS=Aedes aegypti GN=AAEL010068 PE=4 SV=1
CCG011996.1	1068	1	167	0.04023383	5.466322304	7.086017496	Up	0.000587	0.00091768	AAEL008570-PA OS=Aedes aegypti GN=AAEL008570 PE=3 SV=1
CCG015980.1	333	25	167	3.22595605	17.53162829	2.442161307	Up	9.32E-12	2.08E-11	AAEL003906-PA OS=Aedes aegypti GN=AAEL003906 PE=4 SV=1
CCG011934.1	403	32	167	3.41198885	14.48643231	2.086017496	Up	1.77E-13	5.17E-13	-
CCG007295.1	1570	39	167	1.06740105	3.71849186	1.800615277	Up	7.93E-14	2.55E-13	AAEL013462-PA OS=Aedes aegypti GN=AAEL013462 PE=4 SV=1
CCG000184.1	1649	40	167	1.04232225	3.540347011	1.764089401	Up	8.28E-14	2.66E-13	Quinone oxidoreductase OS=Culex quinquefasciatus GN=CpipJ_CPIJ019133 PE=4 SV=1
CCG027725.1	375	41	167	4.69802432	15.56808592	1.728465492	Up	1.35E-13	4.07E-13	AAEL006792-PA OS=Aedes aegypti GN=AAEL006792 PE=4 SV=1
CCG001937.1	3270	387	167	5.08540896	1.785330954	-1.51017226	Down	1.67E-32	7.11E-32	Nfr3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017533 PE=4 SV=1

CCG014444.1	1983	465	167	10.0761102	2.944040454	-1.77506941	Down	1.67E-48	7.74E-48	AAEL014970-PA (Fragment) OS=Aedes aegypti GN=AAEL014970 PE=4 SV=1
CCG017353.1	270	1	168	0.15914717	21.75181666	7.094630627	Up	0.000587	0.0009182	AAEL000323-PA OS=Aedes aegypti GN=AAEL000323 PE=4 SV=1
CCG023510.1	1938	14	168	0.31041088	3.030438853	3.287275705	Up	3.07E-08	6.02E-08	AAEL000199-PA OS=Aedes aegypti GN=AAEL000199 PE=3 SV=1
CCG021244.2	663	22	168	1.42584338	8.858205879	2.635199008	Up	8.38E-11	1.80E-10	AAEL001871-PA OS=Aedes aegypti GN=AAEL001871 PE=4 SV=1
CCG020384.3	1611	32	168	0.8535267	3.645555865	2.094630627	Up	1.77E-13	5.16E-13	AAEL002222-PA OS=Aedes aegypti GN=AAEL002222 PE=4 SV=1
CCG021086.2	306	43	168	6.03823068	19.1927794	1.668365872	Up	2.00E-13	5.77E-13	AAEL001642-PA OS=Aedes aegypti GN=AAEL001642 PE=4 SV=1
CCG027971.1	1566	44	168	1.20732332	3.750313217	1.635199008	Up	2.92E-13	8.09E-13	AAEL013489-PA OS=Aedes aegypti GN=AAEL013489 PE=4 SV=1
CCG002779.1	342	46	168	5.77955495	17.17248684	1.571068671	Up	1.19E-12	2.90E-12	AAEL014732-PA OS=Aedes aegypti GN=AAEL014732 PE=4 SV=2
CCG013329.1	318	54	168	7.29674739	18.46852358	1.339743124	Up	2.66E-10	5.60E-10	AAEL012697-PA OS=Aedes aegypti GN=AAEL012697 PE=4 SV=1
CCG007676.1	948	60	168	2.71960346	6.195137656	1.187740031	Up	8.71E-09	1.74E-08	AAEL000264-PA OS=Aedes aegypti GN=AAEL000264 PE=4 SV=1
CCG023658.1	819	61	168	3.200432	7.170928569	1.163893289	Up	1.49E-08	2.95E-08	AAEL002318-PA OS=Aedes aegypti GN=AAEL002318 PE=4 SV=1
CCG012589.1	1240	62	168	2.14848673	4.736282659	1.140434316	Up	2.52E-08	4.96E-08	AAEL003692-PA OS=Aedes aegypti GN=AAEL003692 PE=4 SV=1
CCG008895.1	972	66	168	2.91769803	6.042171294	1.050236507	Up	1.85E-07	3.50E-07	AAEL001185-PA OS=Aedes aegypti GN=AAEL001185 PE=4 SV=1
CCG019087.1	495	68	168	5.90291304	11.86462727	1.007167785	Up	4.69E-07	8.76E-07	AAEL005495-PA (Fragment) OS=Aedes aegypti GN=AAEL005495 PE=4 SV=1
CCG004238.1	2898	295	168	4.37407581	2.026566769	-1.109940518	Down	3.78E-16	1.42E-15	AAEL007004-PA (Fragment) OS=Aedes aegypti GN=GPRBOS1 PE=4 SV=2
CCG027784.1	930	312	168	14.4156529	6.315043546	-1.190771592	Down	7.60E-19	2.93E-18	AAEL011624-PA OS=Aedes aegypti GN=AAEL011624 PE=3 SV=1

CCG024698.1	1167	335	168	12.3349281	5.032553983	-1.293386659	Down	1.03E-22	4.13E-22	AAEL014372-PA OS=Aedes aegypti GN=AAEL014372 PE=4 SV=1
CCG011123.1	2658	433	168	6.99996053	2.209552482	-1.663592588	Down	1.95E-41	8.76E-41	-
CCG015610.1	1248	634	168	21.8291761	4.705921873	-2.213708404	Down	2.06E-87	1.06E-86	AAEL012370-PA OS=Aedes aegypti GN=AAEL012370 PE=3 SV=1
CCG027943.1	3381	1015	168	12.8998168	1.737057231	-2.892633385	Down	5.37E-189	3.18E-188	Tyrosine kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ000270 PE=3 SV=1
CCG018027.1	1547	1315	168	36.5256632	3.796373948	-3.266216458	Down	7.72E-276	4.81E-275	AAEL001570-PA (Fragment) OS=Aedes aegypti GN=AAEL001570 PE=4 SV=1
CCG027352.3	1191	9	169	0.32470832	4.960494353	3.933267639	Up	1.09E-06	2.01E-06	AAEL005905-PA OS=Aedes aegypti GN=AAEL005905 PE=4 SV=1
CCG006889.1	1083	11	169	0.43644236	5.45516969	3.643761021	Up	2.67E-07	5.04E-07	AAEL015285-PA OS=Aedes aegypti GN=AAEL015285 PE=4 SV=1
CCG008112.1	389	25	169	2.76155107	15.18752898	2.45933645	Up	9.32E-12	2.08E-11	-
CCG008233.1	315	29	169	3.95594382	18.75539293	2.245211645	Up	6.66E-13	1.69E-12	AAEL013332-PA OS=Aedes aegypti GN=AAEL013332 PE=4 SV=1
CCG003658.1	648	38	169	2.51983012	9.117204899	1.855265127	Up	4.37E-14	1.48E-13	AAEL011057-PA OS=Aedes aegypti GN=AAEL011057 PE=4 SV=1
CCG004338.1	2148	44	169	0.88019941	2.750441701	1.643761021	Up	2.14E-13	6.15E-13	AAEL009522-PA OS=Aedes aegypti GN=AAEL009522 PE=4 SV=1
CCG000686.1	1794	276	169	6.6107284	3.293171	-1.005331817	Down	3.80E-13	1.02E-12	AAEL006581-PA OS=Aedes aegypti GN=AAEL006581 PE=4 SV=1
CCG002396.1	292	346	169	50.9161924	20.23270128	-1.331435588	Down	2.06E-24	8.32E-24	-
CCG022625.1	2355	617	169	11.257888	2.508683131	-2.165934039	Down	7.82E-83	3.99E-82	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005715 PE=4 SV=1
CCG005597.1	261	0	170	0.001	22.76975882	14.47483139	Up	4.44E-16	1.66E-15	-
CCG015386.1	642	0	170	0.001	9.256864566	13.1763079	Up	4.44E-16	1.63E-15	AAEL008029-PA OS=Aedes aegypti GN=AAEL008029 PE=3 SV=1
CCG003975.1	396	1	170	0.10850943	15.00734104	7.11170414	Up	0.000587	0.00091734	-
CCG024075.1	279	2	170	0.30802677	21.30074212	6.11170414	Up	9.36E-05	0.00015334	-
CCG004312.1	591	3	170	0.21812048	10.05568029	5.526741639	Up	5.28E-05	8.84E-05	AAEL010150-PA OS=Aedes aegypti GN=AAEL010150 PE=4 SV=1

CCG028618.1	1080	15	170	0.59680187	5.502691714	3.204813544	Up	1.48E-08	2.92E-08	AAEL015285-PA OS=Aedes aegypti GN=AAEL015285 PE=4 SV=1
CCG000980.1	723	42	170	2.49616716	8.219788453	1.719386717	Up	1.01E-13	3.16E-13	AAEL012417-PA OS=Aedes aegypti GN=AAEL012417 PE=4 SV=1
CCG020141.1	1236	61	170	2.1206746	4.808177226	1.180966802	Up	8.29E-09	1.66E-08	AAEL002159-PA OS=Aedes aegypti GN=AAEL002159 PE=4 SV=1
CCG017776.1	1524	66	170	1.86089402	3.899545309	1.067310021	Up	1.07E-07	2.05E-07	AAEL015415-PA OS=Aedes aegypti GN=AAEL015415 PE=4 SV=1
CCG025703.1	777	67	170	3.70524095	7.648529024	1.045614949	Up	1.73E-07	3.28E-07	AAEL012823-PA OS=Aedes aegypti GN=AAEL012823 PE=3 SV=1
CCG019223.1	825	277	170	14.4274139	7.203523699	-1.002038026	Down	3.99E-13	1.06E-12	AAEL001946-PA OS=Aedes aegypti GN=AAEL001946 PE=4 SV=1
CCG015507.2	1122	652	170	24.9699349	5.296708602	-2.237024014	Down	5.14E-91	2.68E-90	AAEL010027-PA OS=Aedes aegypti GN=AAEL010027 PE=4 SV=1
CCG009694.1	561	3310	170	253.529094	10.5934172	-4.580911362	Down	0	0	AAEL001683-PA OS=Aedes aegypti GN=AAEL001683 PE=4 SV=1
CCG014323.1	582	6	171	0.44298695	10.27124627	4.535203218	Up	8.38E-06	1.47E-05	AAEL009773-PA OS=Aedes aegypti GN=AAEL009773 PE=4 SV=1
CCG015543.1	792	8	171	0.43403772	7.547809758	4.120165719	Up	2.19E-06	3.97E-06	AAEL006919-PA OS=Aedes aegypti GN=AAEL006919 PE=3 SV=1
CCG008725.1	1632	32	171	0.84254382	3.662907677	2.120165719	Up	1.77E-13	5.17E-13	AAEL013070-PA OS=Aedes aegypti GN=AAEL013070 PE=4 SV=1
CCG011187.2	914	61	171	2.86778316	6.540334057	1.189428381	Up	6.16E-09	1.24E-08	AAEL012608-PA OS=Aedes aegypti GN=AAEL012608 PE=4 SV=1
CCG019956.2	2880	62	171	0.9250429	2.075647683	1.165969408	Up	1.06E-08	2.11E-08	AAEL011911-PA OS=Aedes aegypti GN=AAEL011911 PE=4 SV=1
CCG005811.1	1984	321	171	6.95226049	3.01303696	-1.206263768	Down	1.03E-19	3.99E-19	AAEL007561-PA OS=Aedes aegypti GN=AAEL007561 PE=4 SV=1
CCG024066.1	1948	403	171	8.88952929	3.068719368	-1.53447031	Down	1.49E-34	6.44E-34	AAEL012920-PA (Fragment) OS=Aedes aegypti GN=GPRALS3 PE=3 SV=2
CCG024846.1	741	619	171	35.8950954	8.06729464	-2.153629881	Down	1.67E-82	8.52E-82	AAEL001294-PA OS=Aedes aegypti GN=AAEL001294 PE=4 SV=1

CCG005100.1	405	902	171	95.7004954	14.7601613	-2.696817905	Down	6.07E-156	3.47E-155	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004234 PE=4 SV=1
CCG002770.1	1131	10	172	0.37992692	5.316378077	3.806649864	Up	5.43E-07	1.01E-06	AAEL011659-PA OS=Aedes aegypti GN=AAEL011659 PE=4 SV=1
CCG018302.2	1377	20	172	0.62410653	4.366611187	2.806649864	Up	3.69E-10	7.72E-10	AAEL002129-PA OS=Aedes aegypti GN=AAEL002129 PE=4 SV=1
CCG007920.2	2226	20	172	0.38607129	2.701178619	2.806649864	Up	3.69E-10	7.71E-10	AAEL006864-PA OS=Aedes aegypti GN=AAEL006864 PE=4 SV=1
CCG001288.1	909	46	172	2.17448602	6.614767442	1.605016002	Up	3.07E-13	8.46E-13	AAEL009957-PA OS=Aedes aegypti GN=AAEL009957 PE=4 SV=1
CCG024695.1	13347	674	172	2.16989594	0.450500008	-2.268026823	Down	1.25E-95	6.55E-95	AAEL009216-PA (Fragment) OS=Aedes aegypti GN=AAEL009216 PE=4 SV=1
CCG006472.1	4413	4211	172	41.0028444	1.362525177	-4.911369202	Down	0	0	r-box/in protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011369 PE=4 SV=1
CCG012363.1	426	15880	172	1601.7826	14.1146094	-6.826345334	Down	0	0	AAEL004249-PA OS=Aedes aegypti GN=AAEL004249 PE=4 SV=1
CCG020442.1	544	0	173	0.001	11.11724611	13.44051184	Up	4.44E-16	1.63E-15	Histone H2B OS=Aedes aegypti GN=AAEL015682 PE=3 SV=1
CCG022130.1	1578	0	173	0.001	3.832561395	11.90409319	Up	4.44E-16	1.63E-15	Alkaline phosphatase OS=Culex quinquefasciatus GN=CpipJ_CPIJ001263 PE=3 SV=1
CCG008249.1	426	2	173	0.20173584	14.19667108	6.136941431	Up	9.36E-05	0.00015324	-
CCG006249.1	1596	10	173	0.26923393	3.789337019	3.815013337	Up	5.43E-07	1.01E-06	AAEL003447-PA OS=Aedes aegypti GN=AAEL003447 PE=4 SV=1
CCG028299.1	857	20	173	1.00279427	7.056921682	2.815013337	Up	3.69E-10	7.71E-10	-
CCG014227.1	417	24	173	2.47307825	14.50307406	2.551978931	Up	1.93E-11	4.23E-11	AAEL003874-PA OS=Aedes aegypti GN=AAEL003874 PE=4 SV=1
CCG018779.1	1008	26	173	1.10834633	5.999783613	2.436501713	Up	4.60E-12	1.05E-11	ETS transcription factor E74B OS=Aedes aegypti GN=E74B PE=2 SV=1
CCG016758.1	762	26	173	1.46615892	7.936721629	2.436501713	Up	4.60E-12	1.05E-11	AAEL009164-PA OS=Aedes aegypti GN=AAEL009164 PE=4 SV=1

CCG013159.1	1989	38	173	0.82094013	3.04061432	1.889013918	Up	4.26E-14	1.45E-13	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010973 PE=4 SV=1
CCG022852.1	540	42	173	3.34209047	11.19959608	1.744624009	Up	8.44E-14	2.70E-13	AAEL005225-PA OS=Aedes aegypti GN=AAEL005225 PE=4 SV=1
CCG023259.1	1203	0	174	0.001	5.056309359	12.30386902	Up	4.44E-16	1.63E-15	AAEL007000-PA OS=Aedes aegypti GN=AAEL007000 PE=4 SV=1
CCG001260.1	1188	15	174	0.54254715	5.120151648	3.238366104	Up	1.48E-08	2.93E-08	AAEL004244-PA OS=Aedes aegypti GN=AAEL004244 PE=4 SV=1
CCG000260.1	1164	35	174	1.29204528	5.225721786	2.015973683	Up	6.39E-14	2.11E-13	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG023620.1	1954	39	174	0.85763544	3.112968351	1.859854481	Up	7.35E-14	2.38E-13	AAEL014628-PA OS=Aedes aegypti GN=AAEL014628 PE=4 SV=1
CCG025074.1	1053	56	174	2.28519006	5.776581347	1.337901778	Up	1.33E-10	2.83E-10	AAEL009581-PA OS=Aedes aegypti GN=AAEL009581 PE=4 SV=1
CCG026410.1	498	349	174	30.1133281	12.21433767	-1.301826527	Down	8.08E-24	3.25E-23	Mitochondrial phosphate carrier protein OS=Aedes aegypti PE=2 SV=1
CCG026490.1	497	388	174	33.5457888	12.2389138	-1.454656143	Down	6.83E-31	2.89E-30	-
CCG003442.1	693	666	174	41.2955891	8.777402826	-2.234121667	Down	8.81E-93	4.60E-92	AAEL012858-PA OS=Aedes aegypti GN=AAEL012858 PE=4 SV=1
CCG026797.1	373	763	174	87.8978754	16.30761437	-2.430282547	Down	2.17E-117	1.18E-116	AAEL003837-PA OS=Aedes aegypti GN=AAEL003837 PE=4 SV=1
CCG004124.1	1931	802	174	17.8465703	3.15004669	-2.502201727	Down	1.41E-127	7.80E-127	AAEL003476-PA OS=Aedes aegypti GN=AAEL003476 PE=4 SV=1
CCG020305.1	1920	1426	174	31.91398	3.168093832	-3.332501567	Down	1.00E-304	6.32E-304	AAEL002376-PA (Fragment) OS=Aedes aegypti GN=AAEL002376 PE=4 SV=2
CCG026159.1	753	2	175	0.11412944	8.124433513	6.153524316	Up	9.36E-05	0.00015329	AAEL015464-PA OS=Aedes aegypti GN=AAEL015464 PE=3 SV=1
CCG010749.1	1725	2	175	0.04981998	3.546491846	6.153524316	Up	9.36E-05	0.00015351	Putative salivary secreted protein 62k-3 OS=Aedes albopictus PE=2 SV=1

CCG023964.1	882	8	175	0.38974816	6.936166026	4.153524316	Up	2.19E-06	3.96E-06	AAEL002232-PA OS=Aedes aegypti GN=AAEL002232 PE=4 SV=1
CCG025169.1	1247	10	175	0.34458488	4.905932987	3.831596221	Up	5.43E-07	1.01E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016173 PE=4 SV=1
CCG006707.1	841	15	175	0.7664043	7.274314429	3.24663372	Up	1.48E-08	2.93E-08	AAEL008962-PA OS=Aedes aegypti GN=AAEL008962 PE=4 SV=1
CCG010220.1	672	17	175	1.08703198	9.103717909	3.066061474	Up	3.39E-09	6.90E-09	AAEL011868-PA OS=Aedes aegypti GN=AAEL011868 PE=4 SV=1
CCG021339.1	1245	31	175	1.06992913	4.913814004	2.199328005	Up	2.40E-13	6.76E-13	AAEL005268-PA OS=Aedes aegypti GN=AAEL005268 PE=4 SV=1
CCG023963.3	1398	40	175	1.22946308	4.376036077	1.831596221	Up	6.91E-14	2.25E-13	AAEL005084-PA OS=Aedes aegypti GN=AAEL005084 PE=4 SV=1
CCG007467.1	1314	54	175	1.7658795	4.655782675	1.398636813	Up	2.69E-11	5.86E-11	AAEL003781-PA OS=Aedes aegypti GN=AAEL003781 PE=3 SV=1
CCG027672.1	648	55	175	3.64712254	9.440892647	1.372164602	Up	5.12E-11	1.10E-10	AAEL010620-PA OS=Aedes aegypti GN=AAEL010620 PE=4 SV=1
CCG019965.1	914	364	175	17.1126733	6.693324327	-1.354270325	Down	3.07E-26	1.26E-25	AAEL009805-PA OS=Aedes aegypti GN=AAEL009805 PE=4 SV=1
CCG015986.1	1359	408	175	12.900406	4.501617686	-1.518901026	Down	1.85E-34	7.98E-34	AAEL010120-PA OS=Aedes aegypti GN=AAEL010120 PE=4 SV=1
CCG024772.1	416	1153	175	119.096404	14.70600585	-3.017652482	Down	5.19E-224	3.14E-223	AAEL002781-PA OS=Aedes aegypti GN=AAEL002781 PE=3 SV=1
CCG003246.1	1258	2125	175	72.5840112	4.863035322	-3.89972281	Down	0	0	-
CCG026578.1	1083	16	176	0.63482526	5.681123464	3.161744822	Up	7.09E-09	1.42E-08	AAEL003146-PA OS=Aedes aegypti GN=AAEL003146 PE=4 SV=1
CCG003681.1	825	20	176	1.04169054	7.457765711	2.839816728	Up	3.69E-10	7.72E-10	AAEL007259-PA OS=Aedes aegypti GN=AAEL007259 PE=4 SV=1
CCG010626.1	1182	25	176	0.90883533	5.205293327	2.517888633	Up	9.32E-12	2.08E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008082 PE=4 SV=1
CCG000885.1	1275	46	176	1.55028062	4.825613107	1.638182866	Up	9.70E-14	3.05E-13	N-acetylglucosamine-6-phosphate deacetylase OS=Aedes aegypti GN=AAEL002430 PE=3 SV=1

CCG015781.1	870	62	176	3.06221097	7.072019209	1.207548512	Up	2.40E-09	4.90E-09	AAEL007688-PA OS=Aedes aegypti GN=AAEL007688 PE=4 SV=1
CCG026896.1	1092	67	176	2.63642145	5.634301018	1.095655632	Up	3.32E-08	6.49E-08	AAEL001451-PA OS=Aedes aegypti GN=AAEL001451 PE=4 SV=1
CCG020140.1	2054	68	176	1.42256181	2.995451174	1.074281981	Up	5.43E-08	1.05E-07	AAEL008101-PA OS=Aedes aegypti GN=AAEL008101 PE=4 SV=1
CCG016715.1	1692	334	176	8.4822053	3.636321934	-1.22195947	Down	7.64E-21	3.00E-20	AAEL014019-PA OS=Aedes aegypti GN=CYP4J17 PE=3 SV=2
CCG021963.1	1068	756	176	30.4167784	5.760914524	-2.400497602	Down	1.21E-114	6.55E-114	AAEL010773-PA OS=Aedes aegypti GN=CLIFE10 PE=3 SV=2
CCG001990.1	1575	1623	176	44.2792884	3.906448706	-3.502702462	Down	0	0	AAEL003194-PA OS=Aedes aegypti GN=AAEL003194 PE=4 SV=1
CCG009608.1	1134	8	177	0.30313746	5.456450607	4.169918754	Up	2.19E-06	3.97E-06	-
CCG021718.1	360	13	177	1.55168486	17.18781941	3.469479036	Up	6.34E-08	1.23E-07	Uncharacterized protein OS=Anopheles darlingi GN=AND_13030 PE=4 SV=1
CCG023582.1	807	13	177	0.69220142	7.667428734	3.469479036	Up	6.34E-08	1.22E-07	AAEL012656-PA OS=Aedes aegypti GN=AAEL012656 PE=4 SV=1
CCG018311.1	888	14	177	0.67745077	6.968034897	3.362563832	Up	3.07E-08	6.01E-08	AAEL002232-PA OS=Aedes aegypti GN=AAEL002232 PE=4 SV=1
CCG000738.1	967	17	177	0.75541416	6.398774549	3.082455913	Up	3.39E-09	6.90E-09	AAEL007541-PA OS=Aedes aegypti GN=AAEL007541 PE=4 SV=1
CCG010146.1	849	18	177	0.91101911	7.288121306	2.999993752	Up	1.62E-09	3.33E-09	AAEL012511-PA OS=Aedes aegypti GN=AAEL012511 PE=4 SV=1
CCG011301.1	1233	20	177	0.69699488	5.018341434	2.847990659	Up	3.69E-10	7.73E-10	AAEL010930-PA OS=Aedes aegypti GN=AAEL010930 PE=4 SV=1
CCG002990.1	714	26	177	1.56472423	8.666127435	2.469479036	Up	4.60E-12	1.05E-11	AAEL007714-PA OS=Aedes aegypti GN=AAEL007714 PE=4 SV=1
CCG017261.1	1149	26	177	0.97233516	5.385217571	2.469479036	Up	4.60E-12	1.05E-11	AAEL008028-PA OS=Aedes aegypti GN=AAEL008028 PE=4 SV=1
CCG010374.1	981	45	177	1.97108874	6.307456665	1.678065658	Up	0	0	AAEL006067-PA OS=Aedes aegypti GN=AAEL006067 PE=4 SV=1
CCG023390.1	3558	49	177	0.59176981	1.739071104	1.55520891	Up	4.09E-13	1.09E-12	AAEL001494-PA OS=Aedes aegypti GN=AAEL001494 PE=4 SV=1

CCG016649.1	1002	53	177	2.27285023	6.17526446	1.441998299	Up	7.08E-12	1.59E-11	AAEL002846-PA OS=Aedes aegypti GN=AAEL002846 PE=4 SV=1
CCG017878.1	1650	354	177	9.21896124	3.75006969	-1.297686796	Down	5.02E-24	2.02E-23	AAEL004258-PA OS=Aedes aegypti GN=AAEL004258 PE=4 SV=1
CCG017635.1	2196	361	177	7.06378606	2.817675314	-1.325936273	Down	3.00E-25	1.22E-24	AAEL014617-PA OS=Aedes aegypti GN=CYP9J28 PE=3 SV=1
CCG026590.1	876	410	177	20.1114055	7.06348743	-1.509561346	Down	2.56E-34	1.10E-33	AGAP008575-PA OS=Anopheles gambiae GN=AGAP008575 PE=4 SV=4
CCG000884.1	2512	415	177	7.09890122	2.463222527	-1.527048772	Down	2.74E-35	1.19E-34	GK21210 OS=Drosophila willistoni GN=GK21210 PE=4 SV=1
CCG023725.1	1329	617	177	19.9490792	4.65584273	-2.099207925	Down	1.27E-79	6.45E-79	AAEL002688-PA OS=Aedes aegypti GN=AAEL002688 PE=3 SV=1
CCG018575.1	1554	1131	177	31.2733397	3.981734227	-2.97346446	Down	1.73E-216	1.04E-215	AAEL003568-PA OS=Aedes aegypti GN=AAEL003568 PE=4 SV=1
CCG009519.1	3699	1874	177	21.7694735	1.672780478	-3.701986484	Down	0	0	Homeotic protein span-major OS=Culex quinquefasciatus GN=CpipJ_CPIJ015319 PE=4 SV=1
CCG018750.1	1081	17	178	0.67574976	5.756311994	3.090583794	Up	3.39E-09	6.89E-09	AAEL006871-PA OS=Aedes aegypti GN=AAEL006871 PE=4 SV=1
CCG027984.1	825	23	178	1.19794412	7.542513049	2.654484679	Up	4.01E-11	8.68E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012744 PE=4 SV=1
CCG020545.1	1569	26	178	0.71205424	3.965948544	2.477606917	Up	4.60E-12	1.05E-11	AAEL008052-PA OS=Aedes aegypti GN=AAEL008052 PE=4 SV=1
CCG007511.1	834	33	178	1.7002413	7.461119023	2.133652515	Up	1.31E-13	4.01E-13	AAEL001425-PA OS=Aedes aegypti GN=AAEL001425 PE=4 SV=1
CCG004036.1	606	38	178	2.69447181	10.26827272	1.930119121	Up	4.26E-14	1.45E-13	AAEL001382-PA OS=Aedes aegypti GN=AAEL001382 PE=4 SV=1
CCG017374.1	939	40	178	1.83044663	6.626808589	1.85611854	Up	6.86E-14	2.24E-13	AAEL010416-PA OS=Aedes aegypti GN=AAEL010416 PE=4 SV=1
CCG021503.1	873	46	178	2.26415555	7.127804428	1.654484679	Up	6.39E-14	2.11E-13	AAEL010239-PA OS=Aedes aegypti GN=AAEL010239 PE=4 SV=1

CCG007575.1	6042	52	178	0.36981566	1.02988634	1.477606917	Up	2.54E-12	5.98E-12	Cadherin 69D OS=Culex quinquefasciatus GN=CpipJ_CPIJ002664 PE=4 SV=1
CCG005421.1	4449	315	178	3.04236152	1.398645373	-1.121161384	Down	2.13E-17	8.11E-17	AAEL010466-PA (Fragment) OS=Aedes aegypti GN=AAEL010466 PE=4 SV=1
CCG006480.1	1134	361	178	13.6790778	5.487278012	-1.317808392	Down	5.03E-25	2.05E-24	vesicular monoamine transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ007766 PE=4 SV=1
CCG014472.1	2010	394	178	8.4229231	3.095807595	-1.444005185	Down	5.14E-31	2.18E-30	AAEL000915-PA OS=Aedes aegypti GN=AAEL000915 PE=4 SV=1
CCG012224.1	970	738	178	32.6924373	6.415023985	-2.349430371	Down	4.08E-109	2.19E-108	AAEL009295-PA OS=Aedes aegypti GN=AAEL009295 PE=4 SV=1
CCG028503.1	2752	919	178	14.3492682	2.261109471	-2.665874417	Down	7.95E-157	4.55E-156	AAEL000590-PA OS=Aedes aegypti GN=DNR1 PE=4 SV=1
CCG011206.2	1005	18	179	0.76960719	6.226399545	3.01620398	Up	1.62E-09	3.33E-09	AAEL013286-PA (Fragment) OS=Aedes aegypti GN=AAEL013286 PE=4 SV=1
CCG008265.1	804	27	179	1.44301348	7.782999431	2.431241479	Up	2.28E-12	5.41E-12	AAEL009778-PA OS=Aedes aegypti GN=AAEL009778 PE=4 SV=1
CCG007031.1	1888	27	179	0.61450362	3.31437052	2.431241479	Up	2.28E-12	5.41E-12	AAEL003034-PA OS=Aedes aegypti GN=AAEL003034 PE=4 SV=1
CCG001657.1	1254	32	179	1.09651635	4.990057051	2.186128981	Up	1.77E-13	5.17E-13	AAEL010376-PA OS=Aedes aegypti GN=AAEL010376 PE=4 SV=1
CCG009082.1	1131	48	179	1.82364921	5.532742301	1.60116648	Up	1.52E-13	4.52E-13	AAEL006666-PA OS=Aedes aegypti GN=AAEL006666 PE=4 SV=1
CCG020728.1	492	59	179	5.15287468	12.71856005	1.303485932	Up	1.69E-10	3.59E-10	Uncharacterized protein OS=Anopheles darlingi GN=AND_20430 PE=4 SV=1
CCG004931.1	1654	70	179	1.81854983	3.783271791	1.056845964	Up	6.27E-08	1.22E-07	AAEL012038-PA OS=Aedes aegypti GN=AAEL012038 PE=4 SV=1
CCG021702.1	2966	586	179	8.48963738	2.109754397	-2.008627873	Down	1.54E-71	7.65E-71	Uncharacterized protein OS=Anopheles darlingi GN=AND_14085 PE=4 SV=1

CCG002522.1	1620	713	179	18.9119881	3.862673792	-2.291629285	Down	2.40E-102	1.28E-101	AAEL000464-PA OS=Aedes aegypti GN=AAEL000464 PE=4 SV=1
CCG000500.1	723	2	180	0.1188651	8.70330542	6.1941663	Up	9.36E-05	0.0001536	AAEL003987-PA OS=Aedes aegypti GN=AAEL003987 PE=4 SV=1
CCG009407.1	1128	3	180	0.11428121	5.578448421	5.609203799	Up	5.28E-05	8.86E-05	AAEL002291-PA OS=Aedes aegypti GN=AAEL002291 PE=4 SV=1
CCG008621.1	1032	9	180	0.37473606	6.097373856	4.024241299	Up	1.09E-06	2.01E-06	AAEL007257-PA OS=Aedes aegypti GN=AAEL007257 PE=4 SV=1
CCG014722.1	2739	16	180	0.25100977	2.297367586	3.1941663	Up	7.09E-09	1.42E-08	AAEL012906-PA OS=Aedes aegypti GN=AAEL012906 PE=4 SV=1
CCG022262.1	446	32	180	3.08303029	14.10872157	2.1941663	Up	1.77E-13	5.16E-13	Anti-microbial peptide cecropin OS=Anopheles darlingi PE=3 SV=1
CCG000843.2	873	46	180	2.26415555	7.207892118	1.670604344	Up	4.82E-14	1.63E-13	Cleavage and polyadenylation specificity factor subunit 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009192 PE=4 SV=1
CCG017971.1	2310	582	180	10.8261409	2.724021567	-1.990709043	Down	3.07E-70	1.52E-69	AAEL004090-PA (Fragment) OS=Aedes aegypti GN=AAEL004090 PE=4 SV=1
CCG007112.1	1617	618	180	16.4225702	3.891459381	-2.077296728	Down	1.09E-78	5.54E-78	AAEL005206-PA OS=Aedes aegypti GN=AAEL005206 PE=4 SV=1
CCG021358.1	1656	1010	180	26.2073864	3.799812693	-2.785973277	Down	1.06E-180	6.24E-180	AAEL011766-PA OS=Aedes aegypti GN=AAEL011766 PE=4 SV=1
CCG024653.1	1626	26	181	0.68709293	3.891419493	2.501719373	Up	4.60E-12	1.05E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002640 PE=4 SV=1
CCG018472.1	1854	28	181	0.64894961	3.412863051	2.394804169	Up	1.17E-12	2.85E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002893 PE=4 SV=1
CCG005540.1	1461	43	181	1.26468076	4.330902187	1.775894336	Up	8.53E-14	2.71E-13	AAEL001870-PA OS=Aedes aegypti GN=AAEL001870 PE=4 SV=1
CCG007056.1	1195	49	181	1.76193891	5.294935645	1.587449247	Up	8.46E-14	2.70E-13	AAEL006548-PA OS=Aedes aegypti GN=AAEL006548 PE=4 SV=1
CCG024257.1	1959	49	181	1.07479173	3.229937772	1.587449247	Up	8.46E-14	2.70E-13	AAEL004194-PA OS=Aedes aegypti GN=AAEL004194 PE=4 SV=1

CCG013042.1	417	55	181	5.66747099	15.17373644	1.420799377	Up	6.96E-12	1.56E-11	AAEL008223-PA OS=Aedes aegypti GN=AAEL008223 PE=4 SV=1
CCG016953.1	1107	60	181	2.32898291	5.715851938	1.295268495	Up	1.64E-10	3.48E-10	AAEL002853-PA OS=Aedes aegypti GN=AAEL002853 PE=4 SV=1
CCG027649.1	903	65	181	3.09305952	7.007140748	1.179791278	Up	2.80E-09	5.70E-09	AAEL007321-PA OS=Aedes aegypti GN=AAEL007321 PE=4 SV=1
CCG017295.1	4509	70	181	0.66708393	1.403292991	1.072876074	Up	3.63E-08	7.08E-08	AAEL015233-PA OS=Aedes aegypti GN=AAEL015233 PE=4 SV=1
CCG016816.1	1650	521	181	13.5680192	3.834817028	-1.822980471	Down	3.99E-56	1.90E-55	AAEL008650-PA OS=Aedes aegypti GN=AAEL008650 PE=4 SV=1
CCG020116.1	8794	1491	181	7.28540759	0.719518774	-3.339905451	Down	0	0	Retrotransposon ninja DNA (Fragment) OS=Drosophila simulans GN=pol PE=4 SV=1
CCG024120.1	572	0	182	0.001	11.12308806	13.44126975	Up	4.44E-16	1.66E-15	AAEL013741-PA OS=Aedes aegypti GN=AAEL013741 PE=4 SV=1
CCG011502.1	603	0	182	0.001	10.55125435	13.3651269	Up	4.44E-16	1.64E-15	AAEL007855-PA OS=Aedes aegypti GN=AAEL007855 PE=4 SV=1
CCG024177.1	839	11	182	0.56336958	7.583321064	3.750676225	Up	2.67E-07	5.03E-07	AAEL009284-PA OS=Aedes aegypti GN=AAEL009284 PE=4 SV=1
CCG023332.1	690	13	182	0.80957471	9.220878801	3.509668126	Up	6.34E-08	1.23E-07	AAEL011580-PA OS=Aedes aegypti GN=AAEL011580 PE=4 SV=1
CCG015261.1	1119	19	182	0.72960229	5.68579658	2.962180331	Up	7.73E-10	1.61E-09	AAEL004955-PA OS=Aedes aegypti GN=AAEL004955 PE=4 SV=1
CCG028027.1	2208	35	182	0.68113257	2.881524625	2.080824827	Up	6.39E-14	2.11E-13	AAEL000425-PA OS=Aedes aegypti GN=AAEL000425 PE=4 SV=1
CCG019451.1	2595	56	182	0.92728522	2.451794363	1.402752922	Up	9.68E-12	2.16E-11	AAEL003726-PA OS=Aedes aegypti GN=AAEL003726 PE=4 SV=1
CCG013930.1	327	56	182	7.35873131	19.45690022	1.402752922	Up	9.68E-12	2.16E-11	AGAP001351-PA OS=Anopheles gambiae GN=AgaP_AGAP001351 PE=4 SV=3
CCG019415.1	2148	65	182	1.30029458	2.96201414	1.187740031	Up	2.08E-09	4.26E-09	AAEL015386-PA OS=Aedes aegypti GN=AAEL015386 PE=4 SV=1
CCG023885.1	3147	310	182	4.23279877	2.021737011	-1.066016561	Down	6.53E-16	2.39E-15	AAEL001804-PA OS=Aedes aegypti GN=AAEL001804 PE=3 SV=1

CCG019950.1	1178	337	182	12.2927	5.401024085	-1.186496937	Down	4.00E-20	1.56E-19	AAEL014909-PA OS=Aedes aegypti GN=AAEL014909 PE=4 SV=1
CCG021921.1	7231	353	182	2.09767893	0.879879183	-1.253416529	Down	8.93E-23	3.57E-22	Tyrosine kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ003860 PE=4 SV=1
CCG009654.1	3144	698	182	9.53971843	2.023666149	-2.236975382	Down	2.64E-97	1.39E-96	AAEL005321-PA OS=Aedes aegypti GN=AAEL005321 PE=4 SV=1
CCG005408.1	1947	1497	182	33.038363	3.267799883	-3.337750662	Down	0	0	AAEL007191-PA OS=Aedes aegypti GN=AAEL007191 PE=2 SV=1
CCG026615.1	750	0	183	0.001	8.529819532	13.0582995	Up	4.44E-16	1.65E-15	AAEL000499-PA OS=Aedes aegypti GN=SPZ1A PE=4 SV=1
CCG007734.1	669	10	183	0.64229798	9.562577951	3.896084947	Up	5.43E-07	1.01E-06	AAEL004787-PA OS=Aedes aegypti GN=AAEL004787 PE=4 SV=1
CCG028545.1	411	12	183	1.25459079	15.56536411	3.633050541	Up	1.30E-07	2.49E-07	AAEL009177-PA OS=Aedes aegypti GN=AAEL009177 PE=4 SV=1
CCG020097.1	954	18	183	0.81074971	6.705832966	3.048088041	Up	1.62E-09	3.33E-09	AAEL001749-PA OS=Aedes aegypti GN=AAEL001749 PE=4 SV=1
CCG025731.1	1239	59	183	2.04617784	5.163329015	1.335369993	Up	4.72E-11	1.02E-10	Argininosuccinate synthase OS=Culex quinquefasciatus GN=CpipJ_CPIJ012882 PE=3 SV=1
CCG017136.1	290	74	183	10.9646909	22.0598781	1.008559676	Up	1.39E-07	2.65E-07	-
CCG017683.1	3574	309	183	3.71506659	1.789973321	-1.053449986	Down	1.37E-15	5.00E-15	AAEL005918-PA OS=Aedes aegypti GN=AAEL005918 PE=3 SV=1
CCG005321.1	1749	355	183	8.72170142	3.657727072	-1.253662172	Down	6.67E-23	2.67E-22	AGAP013390-PA OS=Anopheles gambiae GN=AgaP_AGAP013390 PE=4 SV=1
CCG016819.1	4293	388	183	3.8835912	1.490185104	-1.3818998	Down	1.08E-28	4.53E-28	AAEL004880-PA (Fragment) OS=Aedes aegypti GN=AAEL004880 PE=4 SV=1
CCG010232.1	917	860	183	40.2987696	6.976406379	-2.530179807	Down	1.94E-138	1.09E-137	AAEL001466-PA OS=Aedes aegypti GN=AAEL001466 PE=4 SV=1
CCG020154.1	2445	1031	183	18.1193441	2.616509059	-2.791815575	Down	7.80E-185	4.59E-184	AAEL010028-PA OS=Aedes aegypti GN=AAEL010028 PE=3 SV=1
CCG018694.1	1436	2191	183	65.5617608	4.45498931	-3.879360727	Down	0	0	-

CCG024592.1	470	2411	183	220.425596	13.61141415	-4.017402894	Down	0	0	Putative salivary odorant binding protein 1 OS=Culex quinquefasciatus PE=2 SV=1
CCG003513.1	1155	6	184	0.2232194	5.569110759	4.640912659	Up	8.38E-06	1.48E-05	AAEL004909-PA OS=Aedes aegypti GN=AAEL004909 PE=4 SV=1
CCG014254.1	1910	19	184	0.42744762	3.367708338	2.977947646	Up	7.73E-10	1.60E-09	AAEL008536-PA OS=Aedes aegypti GN=AAEL008536 PE=4 SV=1
CCG021217.1	541	28	184	2.2239419	11.88969118	2.418520238	Up	1.17E-12	2.86E-12	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 SV=1
CCG000535.1	2556	68	184	1.14316978	2.516558265	1.138412319	Up	5.71E-09	1.15E-08	AAEL013903-PB OS=Aedes aegypti GN=AAEL013903 PE=4 SV=1
CCG000437.1	673	346	184	22.0914237	9.557686369	-1.208753068	Down	3.41E-21	1.34E-20	AAEL013852-PA OS=Aedes aegypti GN=ABDA_AEDAE_b PE=2 SV=2
CCG023060.1	1842	483	184	11.2673083	3.49203199	-1.690004219	Down	4.86E-47	2.24E-46	Transporter OS=Aedes aegypti GN=AAEL003860 PE=3 SV=1
CCG006822.1	2766	506	184	7.86069621	2.325496358	-1.757118415	Down	6.65E-52	3.13E-51	AAEL007535-PA OS=Aedes aegypti GN=AAEL007535 PE=4 SV=1
CCG026080.1	489	14	185	1.23021735	13.22552393	3.426339742	Up	3.07E-08	6.00E-08	AAEL007250-PA OS=Aedes aegypti GN=AAEL007250 PE=4 SV=1
CCG001916.1	957	17	185	0.76330772	6.757869595	3.146231823	Up	3.39E-09	6.89E-09	AAEL011648-PA OS=Aedes aegypti GN=AAEL011648 PE=3 SV=1
CCG011346.1	1306	17	185	0.55933039	4.951976419	3.146231823	Up	3.39E-09	6.88E-09	AGAP005368-PA OS=Anopheles gambiae GN=AGAP005368 PE=4 SV=2
CCG007270.1	698	23	185	1.41590816	9.265445849	2.710132708	Up	4.01E-11	8.68E-11	AAEL012938-PA OS=Aedes aegypti GN=AAEL012938 PE=4 SV=1
CCG022065.1	1666	46	185	1.18643925	3.88192149	1.710132708	Up	3.57E-14	1.24E-13	AAEL004604-PA (Fragment) OS=Aedes aegypti GN=AAEL004604 PE=4 SV=1
CCG027474.1	824	57	185	2.97242096	7.848642236	1.40080465	Up	6.90E-12	1.55E-11	AAEL008486-PA OS=Aedes aegypti GN=AAEL008486 PE=4 SV=1
CCG000811.1	769	65	185	3.63203218	8.40998856	1.211326851	Up	8.52E-10	1.76E-09	-
CCG027238.1	288	72	185	10.7424337	22.45583751	1.063769663	Up	3.19E-08	6.24E-08	AAEL007695-PA OS=Aedes aegypti GN=AAEL007695 PE=4 SV=1

CCG019618.1	1702	74	185	1.86824933	3.799812693	1.024241299	Up	8.19E-08	1.58E-07	AAEL005374-PA OS=Aedes aegypti GN=AAEL003374 PE=4 SV=1
CCG022988.1	7731	335	185	1.86196625	0.836538766	-1.154322621	Down	3.23E-19	1.25E-18	AAEL008384-PA OS=Aedes aegypti GN=AAEL008384 PE=3 SV=1
CCG005288.1	387	4054	185	450.127401	16.71132094	-4.751435709	Down	0	0	AAEL002185-PA OS=Aedes aegypti GN=AAEL002185 PE=4 SV=1
CCG005147.1	1662	1	186	0.02585423	3.912298123	7.241472015	Up	0.000587	0.00091774	AAEL006333-PA OS=Aedes aegypti GN=AAEL006333 PE=3 SV=1
CCG011089.1	1107	1	186	0.03881638	5.873748401	7.241472015	Up	0.000587	0.00091609	FXa-directed anticoagulant OS=Aedes albopictus PE=2 SV=1
CCG020249.1	424	6	186	0.60806228	15.33547047	4.656509514	Up	8.38E-06	1.47E-05	AAEL006972-PA OS=Aedes aegypti GN=AAEL006972 PE=4 SV=1
CCG010067.1	681	14	186	0.88337193	9.548075594	3.434117093	Up	3.07E-08	6.00E-08	AAEL006501-PA OS=Aedes aegypti GN=AAEL006501 PE=4 SV=1
CCG022149.1	742	36	186	2.08478497	8.763125983	2.071547013	Up	6.13E-14	2.03E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005826 PE=4 SV=1
CCG003470.1	916	43	186	2.0171382	7.098514716	1.81520726	Up	8.46E-14	2.71E-13	AGAP005314-PA OS=Anopheles gambiae GN=AGAP005314 PE=4 SV=2
CCG005504.1	1503	47	186	1.34369762	4.326173972	1.686883163	Up	0	0	AAEL007564-PA OS=Aedes aegypti GN=AAEL007564 PE=4 SV=1
CCG014871.1	2519	53	186	0.90408731	2.581278078	1.51355156	Up	2.96E-13	8.19E-13	AAEL014398-PA OS=Aedes aegypti GN=AAEL014398 PE=4 SV=1
CCG020595.1	2921	68	186	1.00032248	2.226032003	1.154009174	Up	3.21E-09	6.53E-09	AAEL010494-PA OS=Aedes aegypti GN=AAEL010494 PE=4 SV=1
CCG012462.1	1545	633	186	17.6050757	4.208569242	-2.064589675	Down	6.37E-80	3.24E-79	AAEL001770-PA OS=Aedes aegypti GN=AAEL001770 PE=4 SV=1
CCG026030.1	1815	658	186	15.5780085	3.582501091	-2.120471759	Down	6.91E-86	3.56E-85	AAEL008559-PA OS=Aedes aegypti GN=AAEL008559 PE=3 SV=1
CCG000332.1	837	0	187	0.001	7.81027211	12.9311571	Up	4.44E-16	1.64E-15	AAEL005972-PA OS=Aedes aegypti GN=AAEL005972 PE=4 SV=1
CCG022376.1	971	5	187	0.22126537	6.732438472	4.927279569	Up	1.59E-05	2.76E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005145 PE=4 SV=1

CCG012771.1	3801	20	187	0.226097	1.719862604	2.927279569	Up	3.69E-10	7.71E-10	AAEL010370-PA OS=Aedes aegypti GN=AAEL010370 PE=4 SV=1
CCG026433.1	928	20	187	0.92607187	7.044394134	2.927279569	Up	3.69E-10	7.72E-10	AAEL012839-PA OS=Aedes aegypti GN=AAEL012839 PE=4 SV=1
CCG007648.1	1050	28	187	1.14585959	6.225902625	2.441852742	Up	1.17E-12	2.85E-12	AAEL001249-PA OS=Aedes aegypti GN=AAEL001249 PE=4 SV=1
CCG022998.1	2020	33	187	0.70198081	3.236236513	2.204813544	Up	1.31E-13	4.02E-13	AAEL005083-PA OS=Aedes aegypti GN=AAEL005083 PE=4 SV=1
CCG017556.1	894	45	187	2.1629061	7.312301741	1.757354567	Up	0	0	AAEL008911-PA OS=Aedes aegypti GN=AAEL008911 PE=4 SV=1
CCG024830.1	1759	49	187	1.19699659	3.716428514	1.63449782	Up	0	0	Kruppel-like zinc finger protein, putative OS=Ixodes scapularis GN=IscW_ISCW018847 PE=4 SV=1
CCG008152.1	506	55	187	4.67062333	12.91936315	1.46784795	Up	8.47E-13	2.12E-12	RAB-interacting factor OS=Aedes aegypti PE=2 SV=1
CCG004313.1	601	65	187	4.64730907	10.87720093	1.226839851	Up	4.66E-10	9.72E-10	AAEL003105-PA (Fragment) OS=Aedes aegypti GN=AAEL003105 PE=4 SV=1
CCG002055.1	1005	68	187	2.90740493	6.504674384	1.161744822	Up	2.40E-09	4.90E-09	AAEL002790-PA OS=Aedes aegypti GN=AAEL002790 PE=4 SV=1
CCG018023.1	520	72	187	5.94965556	12.57153415	1.079282662	Up	1.84E-08	3.63E-08	AGAP007614-PA OS=Anopheles gambiae GN=AGAP007614 PE=4 SV=2
CCG003921.1	1491	75	187	2.16145546	4.384438468	1.020388973	Up	7.64E-08	1.47E-07	Cytochrome P450 OS=Aedes albopictus PE=2 SV=1
CCG024877.1	3780	323	187	3.67175245	1.729417396	-1.086182691	Down	5.55E-17	2.10E-16	AAEL010252-PA OS=Aedes aegypti GN=AAEL010252 PE=4 SV=1
CCG017747.1	1977	336	187	7.30289875	3.306625066	-1.143109759	Down	5.41E-19	2.09E-18	AAEL000330-PA OS=Aedes aegypti GN=AAEL000330 PE=2 SV=1
CCG024706.1	4698	346	187	3.16465053	1.391485261	-1.185420564	Down	1.36E-20	5.34E-20	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015717 PE=4 SV=1
CCG023795.2	1812	468	187	11.0981434	3.607725031	-1.621157056	Down	4.55E-43	2.05E-42	AAEL010527-PA OS=Aedes aegypti GN=AAEL010527 PE=4 SV=1

CCG000631.1	1752	584	187	14.3232449	3.731277258	-1.940616895	Down	3.69E-68	1.82E-67	AAEL007282-PA OS=Aedes aegypti GN=AAEL007282 PE=4 SV=1
CCG023732.1	714	10	188	0.60181701	9.204700327	3.934973961	Up	5.43E-07	1.01E-06	AAEL013634-PA OS=Aedes aegypti GN=AAEL013634 PE=4 SV=1
CCG012902.1	1890	13	188	0.29555902	3.477331234	3.556462337	Up	6.34E-08	1.23E-07	AAEL009031-PA OS=Aedes aegypti GN=AAEL009031 PE=4 SV=1
CCG008144.1	813	14	188	0.73994623	8.083832759	3.449547133	Up	3.07E-08	6.01E-08	AAEL001580-PA OS=Aedes aegypti GN=AAEL001580 PE=4 SV=1
CCG008699.1	1227	25	188	0.87550397	5.356280386	2.613045866	Up	9.32E-12	2.08E-11	AAEL009918-PA OS=Aedes aegypti GN=AAEL009918 PE=4 SV=1
CCG026465.1	2106	32	188	0.65291145	3.120681877	2.256902055	Up	1.77E-13	5.17E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006366 PE=4 SV=1
CCG012980.1	2355	33	188	0.60212367	2.79072443	2.212507936	Up	1.31E-13	4.01E-13	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_11683 PE=4 SV=1
CCG017445.1	834	435	188	22.4122716	7.880283013	-1.507969535	Down	2.96E-36	1.29E-35	Membrane glycoprotein LIG-1 (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG011122.1	1638	607	188	15.9234609	4.012305271	-1.988650651	Down	4.42E-73	2.21E-72	AAEL012481-PA (Fragment) OS=Aedes aegypti GN=HPX6 PE=4 SV=2
CCG000272.1	2832	614	188	9.31617834	2.320676565	-2.00519279	Down	1.05E-74	5.26E-74	AAEL011638-PA (Fragment) OS=Aedes aegypti GN=AAEL011638 PE=4 SV=1
CCG007322.2	1284	12	189	0.4015863	5.145727656	3.679593127	Up	1.30E-07	2.49E-07	tRNA methyltransferase OS=Culex quinquefasciatus GN=CpipJ_CPIJ009249 PE=4 SV=1
CCG011365.2	318	17	189	2.29712418	20.77708903	3.177092787	Up	3.39E-09	6.88E-09	AAEL005884-PA OS=Aedes aegypti GN=AAEL005884 PE=4 SV=1
CCG023173.1	498	24	189	2.07083058	13.26729781	2.679593127	Up	1.93E-11	4.23E-11	AAEL010832-PA OS=Aedes aegypti GN=AAEL010832 PE=4 SV=1
CCG024132.1	810	36	189	1.90976598	8.156931247	2.094630627	Up	6.13E-14	2.03E-13	AAEL004513-PA (Fragment) OS=Aedes aegypti GN=AAEL004513 PE=3 SV=1

CCG016874.1	931	50	189	2.30771937	7.096793029	1.620699438	Up	0	0	AAEL0006859-PA OS=Redes aegypti GN=AAEL006859 PE=4 SV_1
CCG002160.1	1083	70	189	2.7773605	6.100751902	1.135272611	Up	3.83E-09	7.76E-09	AAEL001930-PA OS=Redes aegypti GN=AAEL001930 PE=4 SV_1
CCG009010.1	1081	76	189	3.02099892	6.11203914	1.016628115	Up	7.12E-08	1.37E-07	-
CCG009923.1	1956	1389	189	30.5137839	3.377870302	-3.175275256	Down	1.84E-283	1.15E-282	AAEL003247-PA OS=Redes aegypti GN=AAEL003247 PE=4 SV_2
CCG008458.1	513	0	190	0.001	12.94750992	13.66038704	Up	4.44E-16	1.64E-15	AAEL001949-PA OS=Redes aegypti GN=AAEL001949 PE=4 SV_1
CCG002691.1	593	2	190	0.14492322	11.20079694	6.272168812	Up	9.36E-05	0.00015372	-
CCG005082.1	231	8	190	1.48812934	28.75356098	4.272168812	Up	2.19E-06	3.96E-06	-
CCG004650.1	1202	18	190	0.64347356	5.525850738	3.102243811	Up	1.62E-09	3.32E-09	AAEL001996-PA OS=Redes aegypti GN=AAEL001996 PE=4 SV_1
CCG022994.1	2058	26	190	0.54286351	3.227440518	2.571729094	Up	4.60E-12	1.05E-11	AAEL005067-PA OS=Redes aegypti GN=AAEL005067 PE=4 SV_1
CCG015346.1	1018	28	190	1.18187875	6.52462926	2.46481389	Up	1.17E-12	2.86E-12	AAEL000427-PA OS=Redes aegypti GN=AAEL000427 PE=4 SV_1
CCG000665.1	1761	28	190	0.68322122	3.771761832	2.46481389	Up	1.17E-12	2.85E-12	AAEL009100-PA OS=Redes aegypti GN=AAEL009100 PE=3 SV_1
CCG010928.1	1005	28	190	1.19716674	6.609027449	2.46481389	Up	1.17E-12	2.85E-12	-
CCG005148.1	1341	30	190	0.9612916	4.953074263	2.365278217	Up	3.71E-13	9.93E-13	AAEL009669-PA OS=Redes aegypti GN=AAEL009669 PE=4 SV_1
CCG009596.1	1731	38	190	0.94329862	3.837130322	2.024241299	Up	4.26E-14	1.45E-13	AAEL010596-PA OS=Redes aegypti GN=AAEL010596 PE=3 SV_1
CCG022800.1	1308	40	190	1.31405916	5.078037146	1.950240717	Up	6.86E-14	2.24E-13	AAEL011367-PA OS=Redes aegypti GN=AAEL011367 PE=4 SV_1
CCG007064.1	774	45	190	2.49824038	8.58148913	1.780315716	Up	0	0	AAEL002258-PA OS=Redes aegypti GN=AAEL002258 PE=4 SV_1
CCG026242.1	1197	56	190	2.01027998	5.548932821	1.46481389	Up	6.14E-13	1.57E-12	AAEL005947-PA OS=Redes aegypti GN=AAEL005947 PE=4 SV_1
CCG009645.1	1605	319	190	8.54040208	4.138362982	-1.045243802	Down	7.39E-16	2.70E-15	-
CCG002022.1	537	334	190	26.7260547	12.36885025	-1.11153548	Down	4.04E-18	1.55E-17	AAEL003270-PA OS=Redes aegypti GN=AAEL003270 PE=4 SV_1
CCG026644.1	1140	454	190	17.1125083	5.826379462	-1.554379675	Down	1.99E-39	8.85E-39	AAEL013794-PA OS=Redes aegypti GN=AAEL013794 PE=4 SV_1

CCG015231.1	993	606	190	26.2232217	6.688894851	-1.971005171	Down	4.06E-72	2.02E-71	-
CCG007219.1	1224	633	190	22.2220931	5.426529891	-2.033892877	Down	2.15E-78	1.09E-77	AAEL009107-PA OS=Aedes aegypti GN=AAEL009167 PE=3 SV=1
CCG005409.1	1469	721	190	21.0899787	4.521492571	-2.221686637	Down	1.33E-99	7.07E-99	-
CCG005078.1	934	3	191	0.13801842	7.148855314	5.694779531	Up	5.28E-05	8.86E-05	-
CCG026372.1	1680	14	191	0.35808112	3.974423133	3.47238711	Up	3.07E-08	6.00E-08	AAEL014180-PA OS=Aedes aegypti GN=AAEL014180 PE=4 SV=1
CCG008011.1	777	21	191	1.16134418	8.593347315	2.887424609	Up	1.76E-10	3.73E-10	Aisr1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008260 PE=4 SV=1
CCG013646.1	909	40	191	1.89085741	7.345468497	1.957813937	Up	6.86E-14	2.25E-13	AAEL009957-PA OS=Aedes aegypti GN=AAEL009957 PE=4 SV=1
CCG018115.1	1495	61	191	1.75328014	4.46624138	1.349004694	Up	1.24E-11	2.76E-11	AAEL003244-PA OS=Aedes aegypti GN=AAEL003244 PE=4 SV=1
CCG015580.1	2968	69	191	0.99895946	2.249673472	1.171217575	Up	1.27E-09	2.61E-09	AAEL005690-PA OS=Aedes aegypti GN=AAEL005690 PE=4 SV=1
CCG016324.3	1588	362	191	9.79536771	4.204679385	-1.220103855	Down	2.08E-22	8.28E-22	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014309 PE=4 SV=1
CCG021047.1	1059	701	191	28.4436109	6.305033865	-2.173528602	Down	2.82E-94	1.48E-93	AAEL015296-PA OS=Aedes aegypti GN=AAEL015296 PE=3 SV=1
CCG011482.1	3270	707	191	9.29039828	2.041905463	-2.185824373	Down	9.64E-96	5.07E-95	Tyrosine kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ000270 PE=3 SV=1
CCG018579.1	2418	901	191	16.0114685	2.7613858	-2.535641264	Down	2.62E-145	1.48E-144	AAEL002869-PA OS=Aedes aegypti GN=AAEL002869 PE=4 SV=1
CCG015004.1	1164	1532	191	56.5546679	5.736280811	-3.30145855	Down	0	0	AAEL012849-PA OS=Aedes aegypti GN=AAEL012849 PE=4 SV=1
CCG012108.1	372	2	192	0.23102008	18.04298156	6.287275705	Up	9.36E-05	0.0001534	AGAP006823-PA OS=Anopheles gambiae GN=AGAP006823 PE=4 SV=1
CCG009427.1	3255	6	192	0.07920688	2.062055035	4.702313204	Up	8.38E-06	1.47E-05	AAEL008126-PA OS=Aedes aegypti GN=GPRCIR PE=4 SV=2
CCG000168.1	743	16	192	0.92532403	9.033632759	3.287275705	Up	7.09E-09	1.42E-08	-
CCG007587.1	870	21	192	1.03720049	7.714930046	2.894958282	Up	1.76E-10	3.73E-10	AAEL004125-PA OS=Aedes aegypti GN=AAEL004125 PE=4 SV=1

CCG014508.1	927	43	192	1.99320236	7.240549234	1.86101095	Up	8.46E-14	2.70E-13	AAEL007203-PA OS=Aedes aegypti GN=AAEL007203 PE=4 SV=1
CCG011108.1	2231	51	192	0.98227542	3.008511493	1.614850363	Up	0	0	AAEL003763-PA (Fragment) OS=Aedes aegypti GN=CYP329B1 PE=3 SV=1
CCG001976.1	2520	60	192	1.02308892	2.663487754	1.380385109	Up	4.68E-12	1.07E-11	AAEL007693-PA OS=Aedes aegypti GN=AAEL007693 PE=4 SV=1
CCG000301.1	1203	64	192	2.28600417	5.579375844	1.287275705	Up	5.60E-11	1.21E-10	AAEL000315-PA OS=Aedes aegypti GN=AAEL000315 PE=4 SV=1
CCG023856.1	1844	76	192	1.77098689	3.639907343	1.039348191	Up	3.20E-08	6.27E-08	AAEL008544-PA OS=Aedes aegypti GN=AAEL008544 PE=4 SV=1
CCG024972.1	3128	334	192	4.58820056	2.145776579	-1.096428588	Down	9.37E-18	3.57E-17	-
CCG027669.1	3036	423	192	5.9868899	2.210800112	-1.437238149	Down	5.93E-33	2.54E-32	AAEL007619-PA OS=Aedes aegypti GN=TOLL5A PE=4 SV=1
CCG018927.1	1612	558	192	14.8741389	4.163764975	-1.836845607	Down	1.33E-60	6.44E-60	AAEL006969-PA OS=Aedes aegypti GN=CYP6AG7 PE=3 SV=1
CCG006875.2	1221	680	192	23.9307285	5.497124603	-2.122115232	Down	8.74E-89	4.53E-88	Alanylacylase OS=Culex quinquefasciatus GN=CpipJ_CPIJ013671 PE=4 SV=1
CCG017577.1	3069	1714	192	23.9980857	2.187028068	-3.45587569	Down	0	0	AAEL005392-PA OS=Aedes aegypti GN=AAEL005392 PE=4 SV=1
CCG008719.1	1128	8	193	0.30474989	5.981336363	4.294770241	Up	2.19E-06	3.97E-06	DNA replication complex GINS protein PSF2 OS=Aedes aegypti GN=AAEL006660 PE=3 SV=1
CCG015577.3	1554	8	193	0.22120841	4.341665005	4.294770241	Up	2.19E-06	3.96E-06	AAEL013862-PA OS=Aedes aegypti GN=AAEL013862 PE=3 SV=1
CCG018265.1	1767	17	193	0.41340435	3.818306405	3.2073074	Up	3.39E-09	6.89E-09	AAEL009198-PA OS=Aedes aegypti GN=AAEL009198 PE=4 SV=1
CCG006100.1	915	28	193	1.31492084	7.373713024	2.487415319	Up	1.17E-12	2.85E-12	AAEL002928-PA OS=Aedes aegypti GN=AAEL002928 PE=4 SV=1
CCG018558.1	810	42	193	2.22806031	8.329564712	1.902452818	Up	7.66E-14	2.47E-13	AAEL002219-PA OS=Aedes aegypti GN=AAEL002219 PE=4 SV=1
CCG018989.1	1326	42	193	1.36103232	5.088195639	1.902452818	Up	7.66E-14	2.47E-13	AGAP006454-PA OS=Anopheles gambiae GN=AGAP006454 PE=4 SV=4

CCG027195.1	1206	55	193	1.95964793	5.594483762	1.513410528	Up	4.09E-14	1.40E-13	AAEL003657-PA OS=Aedes aegypti GN=AAEL003657 PE=4 SV=1
CCG019671.1	1068	63	193	2.53473154	6.317366495	1.317490318	Up	2.24E-11	4.90E-11	AAEL008608-PA OS=Aedes aegypti GN=AAEL008608 PE=4 SV=1
CCG009607.1	791	64	193	3.47669155	8.529642752	1.294770241	Up	4.09E-11	8.84E-11	AAEL001738-PA OS=Aedes aegypti GN=AAEL001738 PE=4 SV=1
CCG003901.1	656	73	193	4.78169303	10.28498082	1.104945682	Up	5.68E-09	1.14E-08	-
CCG026608.1	802	480	193	25.7175469	8.41265264	-1.612120355	Down	8.74E-44	3.96E-43	-
CCG015339.1	1174	675	193	24.7057673	5.74697395	-2.103973451	Down	3.45E-87	1.78E-86	AAEL003291-PA OS=Aedes aegypti GN=AAEL003291 PE=4 SV=1
CCG009388.1	861	23	194	1.14785586	7.876777809	2.77866409	Up	4.01E-11	8.66E-11	AAEL003451-PA OS=Aedes aegypti GN=AAEL003451 PE=4 SV=1
CCG015763.2	1554	39	194	1.07839102	4.364160678	2.016823827	Up	7.33E-14	2.38E-13	AAEL014412-PA OS=Aedes aegypti GN=CYP304B2 PE=3 SV=1
CCG019092.1	1806	50	194	1.18963828	3.755208025	1.658369856	Up	0	0	AAEL002312-PA OS=Aedes aegypti GN=AAEL002312 PE=4 SV=1
CCG006010.1	1603	54	194	1.44751445	4.230758387	1.547338544	Up	6.22E-15	2.24E-14	AAEL013897-PA OS=Aedes aegypti GN=AAEL013897 PE=4 SV=1
CCG004684.1	402	60	194	6.41339323	16.87041217	1.39533545	Up	2.38E-12	5.62E-12	AAEL001819-PA OS=Aedes aegypti GN=AAEL001819 PE=4 SV=1
CCG010620.1	2873	71	194	1.06190434	2.360565852	1.152478927	Up	1.53E-09	3.14E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004442 PE=4 SV=1
CCG027558.1	4815	76	194	0.67823465	1.408495471	1.054298533	Up	1.87E-08	3.69E-08	AAEL008117-PA OS=Aedes aegypti GN=AAEL008117 PE=4 SV=1
CCG010292.1	1419	350	194	10.5985956	4.779355669	-1.148985066	Down	7.18E-20	2.80E-19	AAEL005284-PA OS=Aedes aegypti GN=AAEL005284 PE=4 SV=1
CCG004310.1	1722	367	194	9.1578935	3.938388905	-1.217410207	Down	1.27E-22	5.07E-22	AAEL010142-PA OS=Aedes aegypti GN=AAEL010142 PE=4 SV=1
CCG018661.1	2373	489	194	8.85469879	2.857945931	-1.631464609	Down	2.52E-45	1.15E-44	AAEL002511-PA (Fragment) OS=Aedes aegypti GN=AAEL002511 PE=4 SV=1
CCG020993.1	1293	505	194	16.7824563	5.245093344	-1.677913532	Down	1.29E-48	5.95E-48	AAEL001899-PA OS=Aedes aegypti GN=AAEL001899 PE=4 SV=1

CCG023042.1	636	588	194	39.7267358	10.66337373	-1.897446299	Down	1.34E-66	6.57E-66	AAEL001817-PA OS=Aedes aegypti GN=AAEL001817 PE=4 SV=1
CCG028356.1	1200	748	194	26.7844679	5.651588078	-2.244668414	Down	1.38E-104	7.35E-104	AAEL011248-PA OS=Aedes aegypti GN=AAEL011248 PE=4 SV=1
CCG026935.1	444	0	195	0.001	15.35329723	13.9062609	Up	4.44E-16	1.63E-15	AAEL0003185-PA OS=Aedes aegypti GN=AAEL003185 PE=4 SV=1
CCG022213.1	594	0	195	0.001	11.47620197	13.48635764	Up	4.44E-16	1.66E-15	AAEL0006892-PA OS=Aedes aegypti GN=AAEL006892 PE=4 SV=1
CCG007632.1	1008	16	195	0.68205928	6.762761876	3.309643518	Up	7.09E-09	1.42E-08	AAEL007481-PA OS=Aedes aegypti GN=AAEL007481 PE=4 SV=1
CCG016101.1	822	32	195	1.67278772	8.293021862	2.309643518	Up	1.77E-13	5.16E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013674 PE=4 SV=1
CCG024682.1	1184	38	195	1.37909621	5.757486462	2.061716004	Up	4.26E-14	1.45E-13	AAEL007270-PA OS=Aedes aegypti GN=AAEL007270 PE=4 SV=1
CCG004649.2	1878	40	195	0.91522331	3.629853019	1.987715423	Up	6.86E-14	2.24E-13	AAEL001966-PA OS=Aedes aegypti GN=AAEL001966 PE=4 SV=1
CCG001946.1	1074	44	195	1.76039881	6.347173157	1.850211899	Up	3.91E-14	1.34E-13	AAEL0005139-PA OS=Aedes aegypti GN=AAEL005139 PE=4 SV=1
CCG000492.1	1233	50	195	1.74248721	5.528681241	1.665787328	Up	0	0	AAEL001590-PA OS=Aedes aegypti GN=AAEL001590 PE=4 SV=1
CCG020188.1	2814	479	195	7.31432227	2.422481866	-1.594238328	Down	5.12E-43	2.31E-42	AAEL0003276-PA OS=Aedes aegypti GN=AAEL003276 PE=3 SV=1
CCG010102.1	693	520	195	32.2428023	9.836744546	-1.712724295	Down	1.88E-51	8.83E-51	AAEL012858-PA OS=Aedes aegypti GN=AAEL012858 PE=4 SV=1
CCG016765.1	1677	6	196	0.15373787	4.085761626	4.732060547	Up	8.38E-06	1.48E-05	AAEL004909-PA OS=Aedes aegypti GN=AAEL004909 PE=4 SV=1
CCG002197.1	635	8	196	0.54135099	10.79027126	4.317023048	Up	2.19E-06	3.97E-06	-
CCG023383.1	681	12	196	0.75717594	10.06141299	3.732060547	Up	1.30E-07	2.49E-07	-
CCG002721.1	552	15	196	1.16765583	12.41272146	3.410132452	Up	1.48E-08	2.93E-08	-
CCG011525.1	1116	38	196	1.46312716	6.13962567	2.069095534	Up	4.26E-14	1.45E-13	Extracellular signal-regulated kinase OS=Sarcophaga crassipalpis GN=erk PE=2 SV=1
CCG026963.1	1380	39	196	1.21436207	4.965088585	2.031620829	Up	7.33E-14	2.38E-13	AAEL005068-PA OS=Aedes aegypti GN=AAEL005068 PE=4 SV=1

CCG012792.1	342	40	196	5.02569995	20.03456797	1.995094953	Up	6.86E-14	2.25E-13	AAEL007444-PA OS=Aedes aegypti GN=AAEL007444 PE=4 SV=1
CCG006982.1	1035	46	196	1.90976598	6.620118113	1.793461092	Up	3.40E-14	1.18E-13	AAEL007607-PA OS=Aedes aegypti GN=AAEL007607 PE=4 SV=1
CCG025008.1	918	46	196	2.15316753	7.463858657	1.793461092	Up	3.40E-14	1.18E-13	AAEL015041-PA OS=Aedes aegypti GN=AAEL015041 PE=3 SV=1
CCG023474.1	1434	50	196	1.49824737	4.778118722	1.673166858	Up	0	0	DNA ligase 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005161 PE=3 SV=1
CCG016131.1	1449	55	196	1.63101132	4.728655795	1.535663334	Up	0	0	AAEL006815-PA OS=Aedes aegypti GN=CYP9J16 PE=3 SV=1
CCG016333.1	3888	75	196	0.82889149	1.762299961	1.088204357	Up	6.64E-09	1.33E-08	Cainmodulin-binding transcription activator OS=Culex quinquefasciatus GN=CpipJ_CPIJ003688 PE=4 SV=1
CCG012307.1	1647	333	196	8.68786984	4.160183514	-1.062355319	Down	6.83E-17	2.58E-16	AAEL010219-PA OS=Aedes aegypti GN=AAEL010219 PE=4 SV=1
CCG026336.1	2562	381	196	6.39011276	2.674403687	-1.25662414	Down	1.51E-24	6.11E-24	-
CCG016582.1	915	1995	196	93.6881099	7.488330325	-3.645149983	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009981 PE=4 SV=1
CCG018504.1	1035	0	197	0.001	6.653894226	12.69998322	Up	4.44E-16	1.64E-15	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG014759.1	852	1	197	0.05043396	8.083075733	7.324365023	Up	0.000587	0.00091763	AAEL002647-PA OS=Aedes aegypti GN=AAEL002647 PE=4 SV=1
CCG014052.1	411	11	197	1.15004156	16.75615699	3.864933405	Up	2.67E-07	5.03E-07	AAEL011472-PA OS=Aedes aegypti GN=AAEL011472 PE=4 SV=1
CCG015486.1	1683	17	197	0.43403772	4.091967038	3.236902182	Up	3.39E-09	6.89E-09	AAEL005783-PA OS=Aedes aegypti GN=AAEL005783 PE=4 SV=1
CCG018109.1	2946	18	197	0.2625442	2.337671597	3.154440022	Up	1.62E-09	3.33E-09	AAEL009477-PA OS=Aedes aegypti GN=AAEL009477 PE=4 SV=1
CCG011765.1	1348	24	197	0.76503979	5.108887629	2.739402523	Up	1.93E-11	4.23E-11	AAEL012505-PA OS=Aedes aegypti GN=AAEL012505 PE=4 SV=1
CCG026348.1	1752	29	197	0.71125702	3.930810801	2.466384028	Up	6.66E-13	1.69E-12	AAEL001015-PA OS=Aedes aegypti GN=AAEL001015 PE=4 SV=1

CCG016564.1	891	30	197	1.44679241	7.729271071	2.417474428	Up	3.71E-13	9.94E-13	AAEL013901-PA OS=Aedes aegypti GN=AAEL013901 PE=4 SV=1
CCG018808.1	1539	32	197	0.89345777	4.474841146	2.324365023	Up	1.77E-13	5.16E-13	AAEL012391-PA OS=Aedes aegypti GN=AAEL012391 PE=4 SV=1
CCG027408.1	3474	50	197	0.61844753	1.982377814	1.680508833	Up	0	0	AAEL002820-PA OS=Aedes aegypti GN=AAEL002820 PE=4 SV=1
CCG004181.1	1130	59	197	2.24355252	6.094496039	1.441721974	Up	4.50E-13	1.18E-12	AAEL000562-PA OS=Aedes aegypti GN=AAEL000562 PE=4 SV=1
CCG018930.1	2964	437	197	6.33528138	2.323475211	-1.447124446	Down	2.51E-34	1.08E-33	AAEL007004-PA (Fragment) OS=Aedes aegypti GN=GPRBOS1 PE=4 SV=2
CCG023908.1	762	456	197	25.7141719	9.037769717	-1.508524991	Down	6.06E-38	2.68E-37	AAEL001850-PA OS=Aedes aegypti GN=AAEL001850 PE=4 SV=1
CCG022390.1	1603	495	197	13.2688825	4.296182485	-1.626919692	Down	1.11E-45	5.08E-45	AAEL000515-PA OS=Aedes aegypti GN=AAEL000515 PE=4 SV=1
CCG021590.1	753	725	197	41.3719224	9.145790869	-2.177472162	Down	1.16E-97	6.10E-97	AAEL008936-PA OS=Aedes aegypti GN=AAEL008936 PE=4 SV=1
CCG011739.1	918	0	198	0.001	7.54002048	12.88035273	Up	4.44E-16	1.64E-15	AAEL000769-PA OS=Aedes aegypti GN=AAEL000769 PE=4 SV=1
CCG022854.1	1178	19	198	0.69306024	5.87583939	3.08374231	Up	7.73E-10	1.60E-09	AAEL012864-PA OS=Aedes aegypti GN=AAEL012864 PE=4 SV=1
CCG026360.1	720	25	198	1.49200467	9.613526112	2.687813634	Up	9.32E-12	2.08E-11	AAEL011715-PA OS=Aedes aegypti GN=AAEL011715 PE=4 SV=1
CCG016259.1	969	32	198	1.41902116	7.143177297	2.331669824	Up	1.77E-13	5.16E-13	AAEL014018-PA OS=Aedes aegypti GN=AAEL014018 PE=4 SV=1
CCG004855.1	785	35	198	1.91584804	8.817501657	2.202386807	Up	6.39E-14	2.11E-13	AAEL008550-PA OS=Aedes aegypti GN=AAEL008550 PE=4 SV=1
CCG003774.2	946	61	198	2.7707757	7.316848627	1.400932486	Up	1.27E-12	3.10E-12	AAEL011007-PA OS=Aedes aegypti GN=AAEL011007 PE=4 SV=1
CCG023012.1	863	62	198	3.0870493	8.02055481	1.377473514	Up	2.23E-12	5.30E-12	-
CCG007923.1	1260	65	198	2.21669266	5.493443493	1.309302011	Up	1.54E-11	3.40E-11	AAEL006870-PA OS=Aedes aegypti GN=AAEL006870 PE=4 SV=1
CCG008599.1	484	65	198	5.77072882	14.30111323	1.309302011	Up	1.54E-11	3.40E-11	AAEL004272-PA OS=Aedes aegypti GN=AAEL004272 PE=4 SV=1

CCG000062.1	774	405	198	22.4841635	8.942814988	-1.330108274	Down	2.93E-28	1.22E-27	AAEL008753-PA OS=Aedes aegypti GN=AAEL008753 PE=4 SV=1
CCG000432.1	3228	488	198	6.49604414	2.144280917	-1.599067514	Down	5.57E-44	2.53E-43	AAEL012287-PA OS=Aedes aegypti GN=AAEL012287 PE=3 SV=1
CCG027889.1	4219	575	198	5.85626864	1.640611235	-1.835748322	Down	2.41E-62	1.17E-61	AAEL004488-PA OS=Aedes aegypti GN=AAEL004488 PE=3 SV=1
CCG002612.1	306	658	198	92.3989718	22.62006144	-2.03027395	Down	3.13E-81	1.59E-80	AAEL011992-PA OS=Aedes aegypti GN=AAEL011992 PE=4 SV=1
CCG027565.1	1758	0	199	0.001	3.957165573	11.95025172	Up	4.44E-16	1.65E-15	60 kDa chaperonin OS=Wolbachia endosymbiont of Drosophila ananassae GN=groL PE=3 SV=1
CCG005093.1	399	7	199	0.75385499	17.43533102	4.531582902	Up	4.31E-06	7.70E-06	-
CCG004078.1	1023	8	199	0.33602921	6.800290399	4.338937824	Up	2.19E-06	3.96E-06	AAEL015060-PA (Fragment) OS=Aedes aegypti GN=AAEL015060 PE=4 SV=1
CCG015151.1	1022	14	199	0.5886265	6.806944303	3.531582902	Up	3.07E-08	6.01E-08	Late trypsin OS=Aedes albopictus PE=3 SV=1
CCG015827.1	1185	15	199	0.54392069	5.870630445	3.432047229	Up	1.48E-08	2.93E-08	AAEL002244-PA OS=Aedes aegypti GN=AAEL002244 PE=4 SV=1
CCG005097.1	1314	20	199	0.65402945	5.294290013	3.017009729	Up	3.69E-10	7.72E-10	AAEL012582-PA OS=Aedes aegypti GN=AAEL012582 PE=4 SV=1
CCG013206.1	654	29	199	1.90538579	10.63715149	2.480956829	Up	6.66E-13	1.69E-12	AAEL001631-PA OS=Aedes aegypti GN=AAEL001631 PE=4 SV=1
CCG005106.1	606	45	199	3.19082188	11.47969815	1.847084728	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003834 PE=4 SV=1
CCG002580.1	1155	48	199	1.7857552	6.023114353	1.753975324	Up	5.35E-14	1.80E-13	AAEL013357-PA OS=Aedes aegypti GN=AAEL013357 PE=4 SV=1
CCG009212.1	678	53	199	3.35899105	10.26061516	1.61101737	Up	0	0	AAEL012258-PA OS=Aedes aegypti GN=AAEL012258 PE=4 SV=1
CCG008147.1	731	59	199	3.46814547	9.516685469	1.456294775	Up	2.20E-13	6.31E-13	Inhibitor of growth protein 4 OS=Harpegnathos saltator GN=EAI_01644 PE=4 SV=1
CCG018381.1	621	344	199	23.8028804	11.20241075	-1.08732693	Down	5.15E-18	1.97E-17	AAEL001509-PA OS=Aedes aegypti GN=AAEL001509 PE=4 SV=1

CCG011234.1	1341	385	199	12.3365756	5.18769357	-1.249776811	Down	1.37E-24	5.55E-24	AAEL009020-PA OS=Aedes aegypti GN=GSTT3 PE=3 SV=2
CCG015891.1	1524	438	199	12.3495694	4.564761862	-1.435849235	Down	5.14E-34	2.21E-33	AAEL015201-PA OS=Aedes aegypti GN=AAEL015201 PE=4 SV=1
CCG021413.1	1461	453	199	13.3232647	4.761599642	-1.484429416	Down	7.58E-37	3.32E-36	AAEL009920-PA OS=Aedes aegypti GN=AAEL009920 PE=4 SV=1
CCG001668.1	6303	463	199	3.1564314	1.103712054	-1.515930559	Down	9.03E-39	4.01E-38	AAEL005529-PA OS=Aedes aegypti GN=AAEL005529 PE=4 SV=1
CCG001039.1	3275	1253	199	16.4400237	2.124182314	-2.952232875	Down	7.24E-238	4.43E-237	AAEL001627-PA OS=Aedes aegypti GN=UAP PE=2 SV=1
CCG014931.1	2274	2878	199	54.3829799	3.059233543	-4.151913052	Down	0	0	AAEL010921-PA OS=Aedes aegypti GN=AAEL010921 PE=4 SV=1
CCG002167.1	1236	0	200	0.001	5.656679089	12.46573961	Up	4.44E-16	1.65E-15	AAEL005842-PA OS=Aedes aegypti GN=AAEL005842 PE=4 SV=1
CCG014703.1	858	20	200	1.00162552	8.148782464	3.024241299	Up	3.69E-10	7.73E-10	AAEL007801-PA OS=Aedes aegypti GN=AAEL007801 PE=4 SV=1
CCG028403.1	2081	21	200	0.43362058	3.359757499	2.953851971	Up	1.76E-10	3.72E-10	-
CCG015613.1	975	24	200	1.05771654	7.170928569	2.761206893	Up	1.93E-11	4.23E-11	AAEL012361-PA OS=Aedes aegypti GN=AAEL012361 PE=4 SV=1
CCG024625.1	1440	30	200	0.8952028	4.855316218	2.439278798	Up	3.71E-13	9.94E-13	AAEL002292-PA OS=Aedes aegypti GN=AAEL002292 PE=4 SV=1
CCG016376.1	999	35	200	1.50544616	6.998654008	2.216886377	Up	6.39E-14	2.11E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013990 PE=4 SV=1
CCG009063.1	756	44	200	2.50088402	9.248221368	1.886737775	Up	3.91E-14	1.34E-13	AAEL007247-PA OS=Aedes aegypti GN=AAEL007247 PE=4 SV=1
CCG013131.1	1836	46	200	1.07658376	3.808091152	1.822607438	Up	3.40E-14	1.18E-13	AAEL006445-PA OS=Aedes aegypti GN=AAEL006445 PE=4 SV=1
CCG006849.1	612	53	200	3.72123519	11.42427345	1.618248939	Up	0	0	AAEL013347-PA OS=Aedes aegypti GN=AAEL013347 PE=3 SV=1
CCG025162.1	959	57	200	2.5539884	7.29056867	1.513279379	Up	4.09E-14	1.40E-13	AAEL001717-PA OS=Aedes aegypti GN=AAEL001717 PE=4 SV=1
CCG023153.3	2361	352	200	6.40633062	2.961311035	-1.113262225	Down	4.67E-19	1.80E-18	AAEL006535-PA OS=Aedes aegypti GN=AAEL006535 PE=4 SV=1

CCG015902.1	1620	371	200	9.84059972	4.315836639	-1.189105983	Down	4.56E-22	1.81E-21	AAEL011756-PA OS=Aedes aegypti GN=AAEL011756 PE=3 SV=1
CCG010222.1	1413	416	200	12.6506791	4.948092961	-1.354270325	Down	9.34E-30	3.93E-29	AAEL012440-PA OS=Aedes aegypti GN=AAEL012440 PE=4 SV=1
CCG003063.1	3126	716	200	9.84207613	2.236613997	-2.137646384	Down	2.95E-94	1.55E-93	AAEL009547-PA OS=Aedes aegypti GN=AAEL009547 PE=3 SV=1
CCG014540.1	2790	770	200	11.8590307	2.505969661	-2.242545242	Down	1.75E-107	9.38E-107	AAEL009795-PA OS=Aedes aegypti GN=AAEL009795 PE=4 SV=1
CCG008490.1	4413	3103	200	30.2141596	1.584331601	-4.253278588	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016210 PE=4 SV=1
CCG010881.1	561	22	201	1.68508763	12.52515799	2.893933276	Up	8.38E-11	1.80E-10	AAEL003435-PA OS=Aedes aegypti GN=AAEL003435 PE=4 SV=1
CCG012440.1	777	35	201	1.93557363	9.043260787	2.224081878	Up	6.39E-14	2.11E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000877 PE=4 SV=1
CCG017312.1	1075	66	201	2.63814185	6.536384773	1.308970776	Up	1.07E-11	2.37E-11	AAEL002324-PA OS=Aedes aegypti GN=AAEL002324 PE=4 SV=1
CCG025376.1	876	77	201	3.77702005	8.021248437	1.086578354	Up	4.45E-09	8.98E-09	AAEL011017-PA OS=Aedes aegypti GN=AAEL011017 PE=4 SV=1
CCG012953.1	661	435	201	28.2781158	10.63027781	-1.411506696	Down	5.73E-33	2.45E-32	-
CCG009147.1	3336	1289	201	16.6031139	2.10629905	-2.978671653	Down	6.64E-247	4.08E-246	AAEL003331-PA OS=Aedes aegypti GN=AAEL003331 PE=4 SV=2
CCG003120.1	603	13	202	0.92637902	11.71073285	3.660084968	Up	6.34E-08	1.23E-07	AAEL003010-PA OS=Aedes aegypti GN=AAEL003010 PE=4 SV=1
CCG012316.1	3127	26	202	0.35727953	2.258257726	2.660084968	Up	4.60E-12	1.05E-11	AGAP005525-PA OS=Anopheles gambiae GN=AGAP005525 PE=4 SV=1
CCG015478.1	543	27	202	2.13661664	13.00473648	2.605637184	Up	2.28E-12	5.41E-12	-
CCG020719.1	1605	51	202	1.36539344	4.399733276	1.688099345	Up	0	0	AAEL003172-PA OS=Aedes aegypti GN=AAEL003172 PE=4 SV=1
CCG010985.1	2001	63	202	1.35287021	3.529021443	1.383244763	Up	1.16E-12	2.83E-12	Testis-specific serine/threonine kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ001596 PE=4 SV=1

CCG004075.1	561	335	202	25.6592889	12.58747221	-1.027492599	Down	3.71E-16	1.39E-15	AAEL015061-PA OS=Aedes aegypti GN=AAEL015061 PE=4 SV=1
CCG015324.1	1613	343	202	9.13739552	4.377911908	-1.06154008	Down	2.47E-17	9.39E-17	AAEL015267-PA OS=Aedes aegypti GN=AAEL013267 PE=4 SV=1
CCG009244.1	921	374	202	17.4491648	7.667287631	-1.186369773	Down	3.71E-22	1.47E-21	AAEL012425-PA OS=Aedes aegypti GN=AAEL012425 PE=4 SV=1
CCG016934.1	3972	388	202	4.19744638	1.777837842	-1.239388156	Down	1.83E-24	7.39E-24	AAEL011178-PA OS=Aedes aegypti GN=AAEL011178 PE=4 SV=1
CCG008690.1	1194	762	202	27.422896	5.914214328	-2.213122501	Down	1.17E-104	6.26E-104	AAEL005611-PA OS=Aedes aegypti GN=AAEL003611 PE=3 SV=1
CCG004589.1	1236	8	203	0.27812126	5.741529276	4.367649121	Up	2.19E-06	3.97E-06	AAEL000721-PA OS=Aedes aegypti GN=AAEL000721 PE=4 SV=1
CCG020036.1	609	13	203	0.91725213	11.65275892	3.667209403	Up	6.34E-08	1.23E-07	AAEL010194-PA OS=Aedes aegypti GN=AAEL010194 PE=4 SV=1
CCG022937.1	1038	18	203	0.74513991	6.836734282	3.19772412	Up	1.62E-09	3.33E-09	AAEL014289-PA OS=Aedes aegypti GN=AAEL014289 PE=4 SV=1
CCG016565.1	894	19	203	0.91322702	7.937953227	3.119721608	Up	7.73E-10	1.61E-09	AAEL013902-PA OS=Aedes aegypti GN=AAEL013902 PE=4 SV=1
CCG021673.1	605	39	203	2.76994983	11.72980196	2.082246902	Up	7.33E-14	2.37E-13	-
CCG010490.1	414	54	203	5.60474799	17.14137726	1.612761619	Up	0	0	-
CCG022047.1	1986	55	203	1.18999769	3.573278039	1.586289407	Up	0	0	AAEL009512-PA OS=Aedes aegypti GN=AAEL009512 PE=4 SV=1
CCG005174.1	984	64	203	2.79477949	7.211920919	1.367649121	Up	1.56E-12	3.77E-12	AAEL004376-PA OS=Aedes aegypti GN=AAEL004376 PE=4 SV=1
CCG008798.1	2437	76	203	1.34004917	2.911994331	1.119721608	Up	1.54E-09	3.16E-09	AAEL005091-PA OS=Aedes aegypti GN=AAEL005091 PE=4 SV=1
CCG012600.1	2820	346	203	5.27217311	2.516500066	-1.066979107	Down	1.32E-17	5.03E-17	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005448 PE=4 SV=1
CCG027790.1	1986	1391	203	30.0961233	3.573278039	-3.074257584	Down	5.89E-275	3.67E-274	AAEL003316-PA OS=Aedes aegypti GN=AAEL003316 PE=4 SV=1
CCG003734.2	1191	22	204	0.79373145	5.987815669	2.915306927	Up	8.38E-11	1.80E-10	AAEL005406-PA OS=Aedes aegypti GN=AAEL005406 PE=4 SV=1

CCG009562.2	1158	37	204	1.37295352	6.158452903	2.16528518	Up	5.17E-14	1.74E-13	AGAP009500-PA (Fragment) OS=Anopheles gambiae GN=AGAP009500 PE=3 SV=4
CCG009882.1	828	41	204	2.1277284	8.61290877	2.017186541	Up	1.02E-13	3.20E-13	AAEL010949-PA OS=Aedes aegypti GN=AAEL010949 PE=4 SV=1
CCG002686.1	516	47	204	3.91390994	13.82071407	1.820149694	Up	0	0	AAEL000772-PA OS=Aedes aegypti GN=AAEL000772 PE=4 SV=1
CCG009323.1	835	49	204	2.52157724	8.540704744	1.760028702	Up	0	0	Survivin OS=Culex quinquefasciatus GN=CpipJ_CPIJ002689 PE=4 SV=1
CCG004434.1	1359	404	204	12.7739314	5.247600045	-1.283472937	Down	9.37E-27	3.87E-26	AGAP007868-PA OS=Anopheles gambiae GN=AGAP007868 PE=4 SV=2
CCG000799.1	1716	8	205	0.2003251	4.176251013	4.381793303	Up	2.19E-06	3.97E-06	AAEL014814-PA OS=Aedes aegypti GN=AAEL014814 PE=4 SV=1
CCG015614.1	1236	9	205	0.31288642	5.798096067	4.211868302	Up	1.09E-06	2.01E-06	AAEL009445-PA OS=Aedes aegypti GN=AAEL009445 PE=4 SV=1
CCG000183.1	1038	10	205	0.41396661	6.90409127	4.059865208	Up	5.43E-07	1.01E-06	AAEL002980-PA OS=Aedes aegypti GN=AAEL002980 PE=4 SV=1
CCG010488.1	1005	19	205	0.81236314	7.130792774	3.13386579	Up	7.73E-10	1.60E-09	AAEL013286-PA (Fragment) OS=Aedes aegypti GN=AAEL013286 PE=4 SV=1
CCG028020.2	774	54	205	2.99788846	9.258975114	1.626905801	Up	0	0	AAEL013197-PA OS=Aedes aegypti GN=AAEL013197 PE=4 SV=1
CCG022241.2	996	60	205	2.58853823	7.195227649	1.474902708	Up	0	0	AAEL012398-PA OS=Aedes aegypti GN=AAEL012398 PE=4 SV=1
CCG024499.1	786	65	205	3.55347678	9.117616715	1.35942549	Up	1.58E-12	3.81E-12	AAEL004867-PA OS=Aedes aegypti GN=AAEL004867 PE=4 SV=1
CCG014276.1	1689	398	205	10.125491	4.243011686	-1.254831317	Down	1.62E-25	6.61E-25	AAEL012981-PA OS=Aedes aegypti GN=AAEL012981 PE=4 SV=1
CCG006595.1	1056	478	205	19.4503155	6.786407896	-1.519073505	Down	4.33E-40	1.93E-39	AAEL000327-PA OS=Aedes aegypti GN=AAEL000327 PE=4 SV=1
CCG025378.1	2667	578	205	9.31252591	2.687081642	-1.793132379	Down	1.07E-60	5.16E-60	AAEL003890-PA OS=Aedes aegypti GN=AAEL003890 PE=3 SV=1

CCG014610.1	2198	644	205	12.5898585	3.260439826	-1.949123575	Down	2.06E-75	1.04E-74	AAEL005678-PA OS=Aedes aegypti GN=AAEL005678 PE=4 SV=1
CCG007425.1	957	649	205	29.1403947	7.488450092	-1.960281365	Down	1.48E-76	7.43E-76	AAEL004096-PA OS=Aedes aegypti GN=AAEL004096 PE=4 SV=1
CCG011921.1	717	1172	205	70.2378368	9.995044265	-2.812963551	Down	1.46E-211	8.79E-211	AAEL006297-PA OS=Aedes aegypti GN=AAEL006297 PE=4 SV=1
CCG004125.1	768	35	206	1.95825613	9.376829447	2.259530714	Up	6.39E-14	2.11E-13	AAEL005705-PA OS=Aedes aegypti GN=AAEL005705 PE=4 SV=1
CCG027192.1	1338	43	206	1.38094065	5.382216005	1.962548976	Up	8.46E-14	2.71E-13	AAEL003700-PA OS=Aedes aegypti GN=AAEL003700 PE=4 SV=1
CCG017083.1	762	53	206	2.98870858	9.450662749	1.660893276	Up	0	0	AAEL006676-PA OS=Aedes aegypti GN=AAEL006676 PE=4 SV=1
CCG027906.1	1194	62	206	2.23125925	6.031327483	1.434617421	Up	1.33E-15	4.84E-15	AAEL015115-PA OS=Aedes aegypti GN=AAEL015115 PE=4 SV=1
CCG019495.2	708	69	206	4.18772837	10.17147601	1.280289274	Up	1.36E-11	3.01E-11	AAEL004291-PA OS=Aedes aegypti GN=AAEL004291 PE=4 SV=1
CCG005189.2	610	542	206	38.1796658	11.80558199	-1.69333531	Down	1.02E-52	4.81E-52	AAEL000711-PA OS=Aedes aegypti GN=AAEL000711 PE=4 SV=1
CCG015450.1	1602	1627	206	43.6402985	4.495259061	-3.279184805	Down	0	0	AAEL008125-PA OS=Aedes aegypti GN=AAEL008125 PE=4 SV=1
CCG008239.3	1404	2285	206	69.932937	5.129205851	-3.769164719	Down	0	0	AGAP000697-PA OS=Anopheles gambiae GN=AgaP_AGAP000697 PE=4 SV=4
CCG027471.1	456	3	207	0.28269562	15.86921775	5.810837661	Up	5.28E-05	8.85E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000199 PE=4 SV=1
CCG026731.1	392	9	207	0.98655003	18.46011044	4.22587516	Up	1.09E-06	2.01E-06	AAEL003821-PA OS=Aedes aegypti GN=AAEL003821 PE=4 SV=1
CCG015442.2	1722	10	207	0.24953388	4.202301563	4.073872066	Up	5.43E-07	1.01E-06	AAEL011836-PA OS=Aedes aegypti GN=AAEL011836 PE=4 SV=1
CCG011888.1	1073	24	207	0.96111242	6.744047802	2.810837661	Up	1.93E-11	4.24E-11	AAEL008623-PA OS=Aedes aegypti GN=AAEL008623 PE=4 SV=1
CCG026118.1	831	36	207	1.86150475	8.708018402	2.22587516	Up	6.13E-14	2.03E-13	AAEL014815-PA (Fragment) OS=Aedes aegypti GN=AAEL014815 PE=4 SV=1

CCG000548.1	1034	48	207	1.99472656	6.99841711	1.810837661	Up	5.35E-14	1.80E-13	AAEL001717-PA OS=Aedes aegypti GN=AAEL001717 PE=4 SV=1
CCG022788.1	912	50	207	2.35579685	7.934608873	1.751943972	Up	0	0	AAEL005279-PA OS=Aedes aegypti GN=AAEL005279 PE=4 SV=1
CCG012808.1	558	54	207	4.15836141	12.968393	1.640912659	Up	0	0	AAEL007902-PA OS=Aedes aegypti GN=AAEL007902 PE=4 SV=1
CCG004744.1	900	67	207	3.19885802	8.040403658	1.329710971	Up	2.82E-12	6.61E-12	AAEL008081-PA OS=Aedes aegypti GN=AAEL008081 PE=4 SV=1
CCG010890.1	1738	511	207	12.6337942	4.163615243	-1.60137932	Down	4.57E-46	2.10E-45	-
CCG014791.1	2337	744	207	13.6797101	3.096432731	-2.14335865	Down	3.35E-98	1.77E-97	AAEL006201-PA OS=Aedes aegypti GN=AAEL006201 PE=4 SV=1
CCG006219.1	2607	853	207	14.0595257	2.775743495	-2.34060177	Down	2.94E-125	1.62E-124	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013200 PE=4 SV=1
CCG022736.1	405	0	208	0.001	17.95388042	14.13200807	Up	4.44E-16	1.63E-15	AAEL007013-PA OS=Aedes aegypti GN=AAEL007013 PE=4 SV=1
CCG002330.1	504	7	208	0.59680187	14.42722533	4.595398	Up	4.31E-06	7.70E-06	-
CCG003580.1	488	12	208	1.05663282	14.90024912	3.817790421	Up	1.30E-07	2.49E-07	AAEL010999-PA OS=Aedes aegypti GN=AAEL010999 PE=4 SV=1
CCG019797.1	1539	30	208	0.83761666	4.724705373	2.495862326	Up	3.71E-13	9.94E-13	AGAP000928-PA OS=Anopheles gambiae GN=AGAP000928 PE=4 SV=3
CCG020423.1	448	45	208	4.31615638	16.2306285	1.910899826	Up	0	0	-
CCG006908.1	897	48	208	2.29938379	8.106267078	1.817790421	Up	5.35E-14	1.80E-13	Methionine aminopeptidase OS=Aedes aegypti GN=AAEL013150 PE=3 SV=1
CCG026154.1	847	49	208	2.48585242	8.58479524	1.788043078	Up	0	0	AAEL003817-PA OS=Aedes aegypti GN=AAEL003817 PE=2 SV=1
CCG023677.2	1245	59	208	2.03631674	5.840418931	1.520109873	Up	0	0	AAEL006413-PA OS=Aedes aegypti GN=AAEL006413 PE=4 SV=1
CCG011619.1	1134	62	208	2.3493153	6.412100149	1.448556612	Up	0	0	AAEL012592-PA OS=Aedes aegypti GN=AAEL012592 PE=4 SV=1
CCG009911.1	1371	64	208	2.00588112	5.303662705	1.402752922	Up	2.16E-13	6.19E-13	AAEL010283-PA OS=Aedes aegypti GN=AAEL010283 PE=4 SV=1

CCG020019.1	585	361	208	26.5163661	12.42960952	-1.093102105	Down	5.60E-19	2.16E-18	AAEL008315-PA OS=Aedes aegypti GN=AAEL008315 PE=4 SV=1
CCG008004.1	2766	398	208	6.18291915	2.62882197	-1.233871699	Down	6.88E-25	2.79E-24	AAEL002141-PA OS=Aedes aegypti GN=AAEL002141 PE=4 SV=1
CCG009446.1	1677	400	208	10.2491913	4.335910297	-1.241103268	Down	3.18E-25	1.30E-24	AAEL010676-PA (Fragment) OS=Aedes aegypti GN=AAEL010676 PE=4 SV=1
CCG001054.1	1471	535	208	15.6280136	4.943114595	-1.660642159	Down	1.15E-50	5.35E-50	AAEL009513-PA (Fragment) OS=Aedes aegypti GN=AAEL009513 PE=4 SV=1
CCG010398.1	3339	23	209	0.29598799	2.188164075	2.88611038	Up	4.01E-11	8.66E-11	AAEL011649-PA OS=Aedes aegypti GN=AAEL011649 PE=4 SV=1
CCG022355.1	1203	26	209	0.92868919	6.07338308	2.709232618	Up	4.60E-12	1.05E-11	AAEL001481-PA OS=Aedes aegypti GN=AAEL001481 PE=4 SV=1
CCG023095.1	1380	29	209	0.90298718	5.294405685	2.551691341	Up	6.66E-13	1.69E-12	AAEL007136-PA OS=Aedes aegypti GN=AAEL007136 PE=4 SV=1
CCG025668.1	618	34	209	2.36403071	11.8224593	2.322209495	Up	8.82E-14	2.79E-13	-
CCG007191.1	327	50	209	6.57029581	22.34336344	1.765816146	Up	0	0	AAEL011433-PA OS=Aedes aegypti GN=AAEL011433 PE=4 SV=1
CCG019607.2	1024	59	209	2.47579526	7.135038912	1.527029287	Up	0	0	Putative integral membrane protein OS=Glossina morsitans morsitans PE=2 SV=1
CCG004808.1	1510	67	209	1.90660412	4.838595924	1.343583145	Up	1.40E-12	3.39E-12	-
CCG026759.1	674	74	209	4.71774534	10.84017781	1.20021897	Up	9.38E-11	2.00E-10	AAEL011974-PA OS=Aedes aegypti GN=AAEL011974 PE=4 SV=1
CCG008251.1	1119	379	209	14.5536456	6.529293874	-1.156381702	Down	1.36E-21	5.38E-21	AAEL005979-PA (Fragment) OS=Aedes aegypti GN=SCR3 PE=4 SV=2
CCG014240.1	3999	420	209	4.51295037	1.827026718	-1.304573182	Down	2.00E-28	8.36E-28	AAEL006849-PA (Fragment) OS=Aedes aegypti GN=AAEL006849 PE=4 SV=1
CCG003601.1	1239	846	209	29.3401093	5.896916744	-2.314841517	Down	1.37E-122	7.53E-122	AAEL012646-PA OS=Aedes aegypti GN=AAEL012646 PE=4 SV=1
CCG028192.1	556	1090	209	84.2392279	13.14079109	-2.680440084	Down	8.80E-187	5.20E-186	-

CCG010909.1	424	9	210	0.91209342	17.31424085	4.24663372	Up	1.09E-06	2.01E-06	Histone H2B OS=Aedes aegypti GN=AAEL015674 PE=3 SV=1
CCG009140.1	1697	24	210	0.60770397	4.3260095	2.831596221	Up	1.93E-11	4.23E-11	AAEL002785-PA OS=Aedes aegypti GN=AAEL002785 PE=4 SV=1
CCG001942.1	399	32	210	3.44619425	18.39909304	2.416558721	Up	1.77E-13	5.16E-13	AAEL002461-PA OS=Aedes aegypti GN=AAEL002461 PE=4 SV=1
CCG018672.1	1182	34	210	1.23601605	6.210861355	2.32909588	Up	8.82E-14	2.80E-13	AAEL009727-PA OS=Aedes aegypti GN=AAEL009727 PE=4 SV=1
CCG011897.1	990	78	210	3.38549424	7.415392043	1.131156503	Up	5.85E-10	1.22E-09	AAEL013769-PA OS=Aedes aegypti GN=AAEL013769 PE=4 SV=1
CCG016220.1	819	414	210	21.7209647	8.963660711	-1.276928236	Down	3.59E-27	1.49E-26	Pyrroline-5-carboxylate reductase OS=Aedes aegypti GN=AAEL007033 PE=3 SV=1
CCG025766.2	1221	491	210	17.2793937	6.012480035	-1.523020493	Down	2.75E-41	1.23E-40	AAEL014422-PA OS=Aedes aegypti GN=AAEL014422 PE=4 SV=1
CCG022487.1	1846	683	210	15.8983363	3.976835386	-1.999183047	Down	1.42E-82	7.26E-82	AAEL012184-PA OS=Aedes aegypti GN=AAEL012184 PE=4 SV=1
CCG010867.1	1794	719	210	17.2214265	4.092105977	-2.073289239	Down	5.04E-91	2.62E-90	Protease III zinc metalloprotease OS=Culex quinquefasciatus GN=CpipJ_CPIJ011546 PE=4 SV=1
CCG021781.1	573	2902	210	217.623333	12.8119339	-4.086273083	Down	0	0	AAEL009066-PA OS=Aedes aegypti GN=AAEL009066 PE=4 SV=1
CCG004528.1	828	1	211	0.05189581	8.908449757	7.423412393	Up	0.000587	0.00091632	AAEL001690-PA OS=Aedes aegypti GN=AAEL001690 PE=3 SV=1
CCG027083.1	1911	18	211	0.40473847	3.859862061	3.253487391	Up	1.62E-09	3.33E-09	AAEL007475-PA OS=Aedes aegypti GN=AAEL007475 PE=4 SV=1
CCG003725.2	1200	25	211	0.8952028	6.146830332	2.779556203	Up	9.32E-12	2.08E-11	AAEL007816-PA OS=Aedes aegypti GN=CYP4D23 PE=3 SV=1
CCG002439.1	948	28	211	1.26914828	7.780797889	2.61605747	Up	1.17E-12	2.86E-12	AAEL011163-PA OS=Aedes aegypti GN=AAEL011163 PE=4 SV=1
CCG018363.1	1173	29	211	1.06233785	6.288317476	2.565431397	Up	6.66E-13	1.69E-12	AAEL008721-PA OS=Aedes aegypti GN=AAEL008721 PE=4 SV=1
CCG000613.1	972	38	211	1.67988674	7.588679423	2.175484879	Up	4.26E-14	1.45E-13	AAEL003384-PA OS=Aedes aegypti GN=AAEL003384 PE=4 SV=1

CCG024084.1	402	40	211	4.27559548	18.34874726	2.101484298	Up	6.86E-14	2.25E-13	AAEL006551-FA OS=Aedes aegypti GN=AAEL006551 PE=4 SV=1
CCG020017.1	618	41	211	2.85074291	11.93559288	2.065860388	Up	1.02E-13	3.19E-13	AAEL008316-FA OS=Aedes aegypti GN=AAEL008316 PE=4 SV=1
CCG005856.1	1935	45	211	0.99929615	3.811987803	1.931559296	Up	0	0	AAEL011904-FA OS=Aedes aegypti GN=AAEL011904 PE=4 SV=1
CCG002485.1	1330	48	211	1.55078741	5.54601233	1.838449892	Up	5.35E-14	1.80E-13	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC011569 PE=4 SV=1
CCG017420.1	462	57	211	5.30146076	15.96579307	1.590522378	Up	0	0	AAEL012071-FA OS=Aedes aegypti GN=AAEL012071 PE=4 SV=1
CCG010238.1	228	65	211	12.2501436	32.35173859	1.401044579	Up	1.73E-13	5.06E-13	AAEL002252-FA OS=Aedes aegypti GN=AAEL002252 PE=4 SV=1
CCG023436.1	639	66	211	4.43818855	11.54334335	1.379018273	Up	1.69E-13	4.96E-13	Exosomal 3'-5' exoribonuclease complex subunit ski4 (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG006939.1	712	74	211	4.46595556	10.3598264	1.213959027	Up	5.16E-11	1.11E-10	AAEL011092-FA OS=Aedes aegypti GN=AAEL011092 PE=4 SV=1
CCG000526.1	2223	82	211	1.58502845	3.318127035	1.065860388	Up	3.24E-09	6.59E-09	Integrin beta (Fragment) OS=Aedes aegypti GN=AAEL001829 PE=3 SV=1
CCG022170.1	1723	38	212	0.94767842	4.301308576	2.182306145	Up	4.26E-14	1.45E-13	AAEL002529-FA OS=Aedes aegypti GN=AAEL002529 PE=4 SV=1
CCG003252.1	652	38	212	2.50437104	11.36680165	2.182306145	Up	4.26E-14	1.45E-13	Defensin D (Fragment) OS=Aedes albopictus GN=DefD PE=2 SV=1
CCG011367.1	552	42	212	3.26943633	13.42600485	2.037916236	Up	7.66E-14	2.47E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013877 PE=4 SV=1
CCG003782.1	1545	44	212	1.22373354	4.796863868	1.97080204	Up	3.91E-14	1.35E-13	-
CCG015927.1	411	45	212	4.70471547	18.03200651	1.938380562	Up	0	0	AAEL010634-FA OS=Aedes aegypti GN=AAEL010634 PE=4 SV=1
CCG001568.1	919	48	212	2.2443387	8.064368526	1.845271158	Up	5.35E-14	1.79E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_13864 PE=4 SV=1

CCG008815.1	1076	75	212	2.99510232	6.887690219	1.201414968	Up	6.63E-11	1.43E-10	Thyroid hormone receptor-associated protein 6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004789 PE=4 SV=1
CCG023114.1	1638	453	212	11.8835713	4.524514454	-1.393133582	Down	1.29E-33	5.52E-33	AAEL007036-FA OS=Aedes aegypti GN=AAEL007038 PE=4 SV=1
CCG005277.1	3426	529	212	6.63484811	2.163209187	-1.616890254	Down	2.80E-48	1.30E-47	AAEL004222-FA OS=Aedes aegypti GN=AAEL004222 PE=3 SV=1
CCG004977.1	1149	679	212	25.3929067	6.450091102	-1.977034106	Down	6.52E-81	3.32E-80	AAEL013984-FA OS=Aedes aegypti GN=AAEL013984 PE=4 SV=1
CCG022450.1	1632	701	212	18.4569755	4.541148698	-2.023036976	Down	5.08E-86	2.61E-85	AAEL000788-FA OS=Aedes aegypti GN=AAEL000788 PE=4 SV=1
CCG007503.1	730	0	213	0.001	10.20015473	13.31630342	Up	4.44E-16	1.64E-15	AAEL012453-FA OS=Aedes aegypti GN=AAEL012453 PE=4 SV=1
CCG009604.1	885	14	213	0.67974721	8.413686952	3.629667902	Up	3.07E-08	6.01E-08	AAEL005577-FA OS=Aedes aegypti GN=AAEL005577 PE=4 SV=1
CCG005256.1	1155	35	213	1.30211317	6.446851041	2.307739807	Up	6.39E-14	2.11E-13	AAEL011765-FA OS=Aedes aegypti GN=AAEL011765 PE=4 SV=1
CCG010103.1	1097	36	213	1.41012803	6.787705517	2.267097823	Up	6.13E-14	2.03E-13	AAEL012855-FA OS=Aedes aegypti GN=AAEL012855 PE=4 SV=1
CCG022931.2	8020	85	213	0.45541489	0.928443012	1.027631888	Up	7.78E-09	1.56E-08	AAEL003987-FA OS=Aedes aegypti GN=AAEL003987 PE=4 SV=1
CCG011649.1	807	362	213	19.2751474	9.226905765	-1.062823063	Down	3.06E-18	1.17E-17	Veli, isoform C OS=Drosophila melanogaster GN=veli PE=4 SV=2
CCG011793.1	904	377	213	17.9199004	8.236850611	-1.121397889	Down	1.60E-20	6.27E-20	AAEL013640-FA OS=Aedes aegypti GN=AAEL013640 PE=3 SV=1
CCG023645.1	928	855	213	39.5895723	8.023828613	-2.302757786	Down	4.37E-123	2.40E-122	AAEL001447-FA OS=Aedes aegypti GN=AAEL001447 PE=4 SV=1
CCG005162.1	1572	865	213	23.6442878	4.736713074	-2.319533498	Down	1.31E-125	7.24E-125	Partial acyl-CoA reductase 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017921 PE=4 SV=1
CCG014885.1	858	1276	213	63.9037079	8.678453325	-2.88038979	Down	4.84E-236	2.96E-235	AAEL000153-FA OS=Aedes aegypti GN=AAEL000153 PE=4 SV=1
CCG010904.1	443	0	214	0.001	16.88729397	14.04365055	Up	4.44E-16	1.63E-15	Histone H2B OS=Aedes aegypti GN=AAEL015674 PE=3 SV=1

CCG017817.1	732	7	214	0.41091276	10.22004266	4.636425268	Up	4.31E-06	7.71E-06	AAEL007174-PA OS=Aedes aegypti GN=AAEL007174 PE=4 SV=1
CCG024401.1	1819	25	214	0.5905681	4.112738444	2.799924	Up	9.32E-12	2.08E-11	AAEL002739-PA OS=Aedes aegypti GN=AAEL002739 PE=4 SV=1
CCG010983.1	1382	35	214	1.08823496	5.413220861	2.314497173	Up	6.39E-14	2.11E-13	AAEL009798-PA OS=Aedes aegypti GN=AAEL009798 PE=4 SV=1
CCG011129.1	1016	54	214	2.28382448	7.363259084	1.688892688	Up	0	0	AAEL003786-PA OS=Aedes aegypti GN=AAEL003786 PE=4 SV=1
CCG011200.2	2628	57	214	0.93199196	2.84667855	1.610890176	Up	0	0	AAEL009432-PA OS=Aedes aegypti GN=SCRBQ3 PE=4 SV=2
CCG013705.1	540	68	214	5.41100362	13.85383561	1.356317349	Up	5.24E-13	1.35E-12	Putative glycoprotein hormone-alpha2 OS=Aedes aegypti GN=gpa2 PE=2 SV=1
CCG025323.1	3069	647	214	9.05878732	2.437625034	-1.893841712	Down	6.45E-73	3.22E-72	Transporter OS=Aedes aegypti GN=AAEL003641 PE=3 SV=1
CCG012433.1	1275	750	214	25.2763145	5.867506846	-2.106966595	Down	7.56E-97	3.98E-96	AAEL000024-PA OS=Aedes aegypti GN=AAEL000024 PE=4 SV=1
CCG010161.1	560	1460	214	112.028237	13.35905577	-3.067972464	Down	6.61E-288	4.14E-287	-
CCG011450.1	5877	6795	214	49.6816993	1.272940485	-5.286477645	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008587 PE=4 SV=1
CCG001799.1	1134	4	215	0.15156873	6.627891981	5.450506053	Up	2.95E-05	5.03E-05	AAEL007686-PA OS=Aedes aegypti GN=AAEL007686 PE=4 SV=1
CCG005284.1	3420	36	215	0.452313	2.197669446	2.280581052	Up	6.13E-14	2.03E-13	AAEL011780-PA OS=Aedes aegypti GN=AAEL011780 PE=3 SV=1
CCG022706.1	1563	43	215	1.18214881	4.808720093	2.024241299	Up	8.46E-14	2.70E-13	AAEL012566-PA OS=Aedes aegypti GN=AAEL012566 PE=4 SV=1
CCG004853.1	1689	62	215	1.5773378	4.449987866	1.496309743	Up	0	0	AAEL008522-PA OS=Aedes aegypti GN=AAEL008522 PE=4 SV=1
CCG015963.1	813	74	215	3.91114436	9.244808741	1.241052688	Up	1.54E-11	3.40E-11	SCO1, mitochondrial OS=Culex quinquefasciatus GN=CpipJ_CPIJ003089 PE=4 SV=1
CCG026085.1	1189	436	215	15.756774	6.321303201	-1.317678271	Down	6.99E-30	2.94E-29	AAEL015598-PA OS=Aedes aegypti GN=AAEL015598 PE=4 SV=1

CCG026933.1	2358	442	215	8.05454737	3.187459502	-1.337396506	Down	6.13E-31	2.60E-30	AAEL007804-PA (Fragment) OS=Aedes aegypti GN=AAEL007804 PE=4 SV=1
CCG025135.1	2607	541	215	8.91700285	2.883018606	-1.62897873	Down	7.45E-50	3.47E-49	AAEL005645-PA OS=Aedes aegypti GN=AAEL005645 PE=3 SV=1
CCG019954.3	1089	1658	215	65.4213223	6.901771815	-3.244722238	Down	0	0	Phosphoserine aminotransferase OS=Aedes aegypti PE=2 SV=1
CCG006975.1	465	33	216	3.04946504	16.2386834	2.412806587	Up	1.31E-13	4.01E-13	AGAP008101-PA (Fragment) OS=Anopheles gambiae GN=AGAP008101 PE=4 SV=4
CCG024927.1	552	68	216	5.2933731	13.67932569	1.369737865	Up	2.44E-13	6.87E-13	AAEL006134-PA OS=Aedes aegypti GN=AAEL006134 PE=4 SV=1
CCG026799.1	2076	87	216	1.80075477	3.637277352	1.01425721	Up	8.83E-09	1.76E-08	Transporter OS=Aedes aegypti GN=AAEL015287 PE=3 SV=1
CCG024554.1	1494	15	217	0.43142304	5.077607804	3.556973841	Up	1.48E-08	2.93E-08	AAEL010046-PA OS=Aedes aegypti GN=AAEL010046 PE=4 SV=1
CCG010635.1	798	22	217	1.18462927	9.50619807	3.004432818	Up	8.38E-11	1.80E-10	AAEL010046-PA OS=Aedes aegypti GN=AAEL010046 PE=4 SV=1
CCG007887.2	642	23	217	1.53941417	11.81611536	2.94030248	Up	4.01E-11	8.68E-11	AAEL000692-PA OS=Aedes aegypti GN=AAEL000692 PE=4 SV=1
CCG014050.1	2022	23	217	0.48877542	3.751704283	2.94030248	Up	4.01E-11	8.67E-11	AAEL011477-PA OS=Aedes aegypti GN=AAEL011477 PE=4 SV=1
CCG021508.1	2991	23	217	0.33042591	2.536257459	2.94030248	Up	4.01E-11	8.68E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_07180 PE=4 SV=1
CCG019604.1	1471	375	217	10.9542151	5.156999361	-1.086882349	Down	1.74E-19	6.75E-19	AAEL007363-PA OS=Aedes aegypti GN=AAEL007363 PE=4 SV=1
CCG002851.1	450	414	217	39.5321558	16.85765791	-1.229622521	Down	1.08E-25	4.42E-25	AAEL003459-PA OS=Aedes aegypti GN=AAEL003459 PE=4 SV=1
CCG009619.1	522	870	217	71.6162244	14.53246372	-2.301007154	Down	4.27E-125	2.36E-124	AAEL009627-PA OS=Aedes aegypti GN=AAEL009627 PE=4 SV=1
CCG018011.1	2274	7	218	0.13227271	3.351321168	4.663142607	Up	4.31E-06	7.71E-06	AAEL001477-PA OS=Aedes aegypti GN=AAEL001477 PE=4 SV=1

CCG016179.1	1019	10	218	0.42168532	7.478807003	4.148569434	Up	5.43E-07	1.01E-06	Pol polyprotein (Fragment) OS=Drosophila buzzatii GN=pol PE=4 SV=1
CCG008689.1	984	19	218	0.82970016	7.74482148	3.222570015	Up	7.73E-10	1.60E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010331 PE=4 SV=1
CCG006043.1	399	20	218	2.15387141	19.10001087	3.148569434	Up	3.69E-10	7.71E-10	AAEL010832-PA OS=Aedes aegypti GN=AAEL010832 PE=4 SV=1
CCG017143.1	1929	39	218	0.86875047	3.950702092	2.18509531	Up	7.33E-14	2.37E-13	AAEL005636-PA OS=Aedes aegypti GN=AAEL005636 PE=4 SV=1
CCG010653.1	1185	42	218	1.52297794	6.4311429	2.078180106	Up	7.66E-14	2.47E-13	AAEL000241-PA OS=Aedes aegypti GN=AAEL000241 PE=4 SV=1
CCG006703.1	1370	47	218	1.47414418	5.562703895	1.915908677	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000155 PE=4 SV=1
CCG001676.1	686	50	218	3.13190486	11.10918999	1.826641339	Up	0	0	5-hydroxyisourate hydrolase OS=Culex quinquefasciatus GN=CpipJ_CPIJ014057 PE=3 SV=1
CCG019359.1	633	66	218	4.48025669	12.03934334	1.426103409	Up	0	0	AAEL005204-PA OS=Aedes aegypti GN=AAEL005204 PE=4 SV=1
CCG000555.1	385	83	218	9.26360512	19.79455672	1.095458097	Up	7.67E-10	1.59E-09	-
CCG019159.1	2148	384	218	7.68174027	3.547907047	-1.114464972	Down	1.11E-20	4.36E-20	AAEL002774-PA OS=Aedes aegypti GN=AAEL002774 PE=4 SV=1
CCG014893.1	709	499	218	30.2424507	10.74880724	-1.492398477	Down	9.94E-41	4.45E-40	-
CCG020988.1	1326	672	218	21.7765171	5.747288338	-1.921819894	Down	3.70E-77	1.87E-76	AAEL009167-PA OS=Aedes aegypti GN=AAEL009167 PE=3 SV=1
CCG010304.1	438	854	218	83.7811721	17.39932497	-2.267594731	Down	1.20E-120	6.57E-120	AAEL012513-PA OS=Aedes aegypti GN=AAEL012513 PE=4 SV=1
CCG015521.1	2190	10	219	0.19620883	3.495827677	4.155172169	Up	5.43E-07	1.01E-06	AAEL004017-PA OS=Aedes aegypti GN=AAEL004017 PE=4 SV=1
CCG014577.1	810	83	219	4.40307157	9.451682238	1.102060832	Up	5.80E-10	1.21E-09	AAEL012610-PA OS=Aedes aegypti GN=AAEL012610 PE=2 SV=1
CCG009393.1	1036	18	220	0.7465784	7.423572288	3.313747916	Up	1.62E-09	3.33E-09	AAEL008576-PA OS=Aedes aegypti GN=AAEL008576 PE=4 SV=1

CCG022397.1	507	27	220	2.28832906	15.16927197	2.728785415	Up	2.28E-12	5.41E-12	AAEL007988-PA OS=Aedes aegypti GN=AAEL007988 PE=4 SV=1
CCG005900.2	2919	33	220	0.48578323	2.63474508	2.439278798	Up	1.31E-13	4.01E-13	AAEL008183-PA OS=Aedes aegypti GN=AAEL008183 PE=4 SV=1
CCG013163.1	963	70	220	3.12344904	7.986314527	1.3543899	Up	2.07E-13	5.93E-13	-
CCG008939.1	1152	72	220	2.68560841	6.6760598	1.313747916	Up	9.54E-13	2.37E-12	AGAP006521-PA OS=Anopheles gambiae GN=AGAP006521 PE=4 SV=4
CCG010433.1	468	76	220	6.97799109	16.43337797	1.235745404	Up	1.05E-11	2.32E-11	AAEL001105-PA OS=Aedes aegypti GN=AAEL001105 PE=4 SV=1
CCG019985.2	1691	624	220	15.8563657	4.548090414	-1.801729302	Down	8.15E-66	3.99E-65	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014431 PE=4 SV=1
CCG000287.1	258	1007	220	167.715204	29.80938329	-2.492175051	Down	5.85E-159	3.36E-158	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015032 PE=4 SV=1
CCG021580.1	699	21	221	1.29093623	11.05261683	3.09789834	Up	1.76E-10	3.73E-10	AAEL006839-PA OS=Aedes aegypti GN=AAEL006839 PE=4 SV=1
CCG012403.1	2491	51	221	0.87974968	3.101476984	1.817790421	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009352 PE=4 SV=1
CCG022330.1	442	55	221	5.34691268	17.47913839	1.70885605	Up	0	0	-
CCG026236.1	339	55	221	6.97149087	22.78990905	1.70885605	Up	0	0	AAEL005501-PA OS=Aedes aegypti GN=AAEL005501 PE=4 SV=1
CCG014384.1	1166	79	221	2.9113285	6.625882647	1.186435015	Up	4.08E-11	8.80E-11	AAEL005258-PA OS=Aedes aegypti GN=AAEL005258 PE=4 SV=1
CCG017241.1	723	80	221	4.75460411	10.68572499	1.168287668	Up	6.97E-11	1.50E-10	AAEL001202-PA OS=Aedes aegypti GN=AAEL001202 PE=4 SV=1
CCG003878.1	1389	85	221	2.6295374	5.562116031	1.080824827	Up	8.93E-10	1.84E-09	AAEL013263-PA OS=Aedes aegypti GN=AAEL013263 PE=4 SV=1
CCG022103.1	630	401	221	27.3505771	12.26314153	-1.157242663	Down	8.85E-23	3.53E-22	AAEL007763-PA OS=Aedes aegypti GN=AAEL007763 PE=4 SV=1
CCG026314.1	873	714	221	35.1436317	8.849689767	-1.989564501	Down	1.07E-85	5.51E-85	AAEL010783-PA OS=Aedes aegypti GN=AAEL010783 PE=3 SV=1

CCG022264.1	11563	1068	221	3.96883824	0.668146603	-2.570480168	Down	1.33E-174	7.72E-174	AAEL014889-PA (Fragment) OS=Aedes aegypti GN=AAEL014889 PE=2 SV=1
CCG022545.1	1062	1155	221	46.732621	7.274744978	-2.683461373	Down	4.72E-198	2.81E-197	AGAP003014-PA OS=Anopheles gambiae GN=AgaP_AGAP003014 PE=4 SV=4
CCG023431.1	1281	24	222	0.80505358	6.058343047	2.911766569	Up	1.93E-11	4.23E-11	AAEL008050-PA OS=Aedes aegypti GN=AAEL008050 PE=4 SV=1
CCG024003.1	1357	29	222	0.91829204	5.71904012	2.638748075	Up	6.66E-13	1.69E-12	AAEL008798-PA OS=Aedes aegypti GN=AAEL008798 PE=4 SV=1
CCG023377.2	2067	30	222	0.62365362	3.754589958	2.589838475	Up	3.71E-13	9.94E-13	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018423 PE=4 SV=1
CCG009033.1	943	36	222	1.64041405	8.229838222	2.326804069	Up	6.13E-14	2.03E-13	AAEL000659-PA OS=Aedes aegypti GN=AAEL000659 PE=4 SV=1
CCG013619.1	1323	62	222	2.01369883	5.866014696	1.54253276	Up	0	0	AAEL015215-PA OS=Aedes aegypti GN=AAEL015215 PE=4 SV=1
CCG013674.1	2043	71	222	1.49331922	3.798696742	1.346981951	Up	1.20E-13	3.69E-13	AAEL004623-PA OS=Aedes aegypti GN=AAEL004623 PE=4 SV=1
CCG012674.1	1044	80	222	3.29269997	7.433656555	1.174800975	Up	5.21E-11	1.12E-10	AAEL001417-PA OS=Aedes aegypti GN=AAEL001417 PE=4 SV=1
CCG022056.1	1163	88	222	3.25136427	6.673033055	1.037297452	Up	2.85E-09	5.81E-09	NAD-dependent protein deacetylase OS=Aedes aegypti GN=AAEL004004 PE=3 SV=1
CCG000647.1	1446	0	223	0.001	5.391214191	12.39639451	Up	4.44E-16	1.64E-15	Angiopoietin like salivary protein OS=Aedes albopictus PE=2 SV=1
CCG003430.2	3218	3	223	0.0400588	2.422528191	5.918250603	Up	5.28E-05	8.84E-05	Drosophila melanogaster SD07741p OS=Plasmodium yoelii yoelii GN=PY02675 PE=4 SV=1
CCG022445.1	1632	10	223	0.26329494	4.776774338	4.181285009	Up	5.43E-07	1.01E-06	AAEL010492-PA OS=Aedes aegypti GN=AAEL010492 PE=4 SV=1
CCG023549.1	1266	12	223	0.40729606	6.157737536	3.918250603	Up	1.30E-07	2.49E-07	AAEL014934-PA OS=Aedes aegypti GN=AAEL014934 PE=4 SV=1
CCG014558.1	2391	16	223	0.28754318	3.260433174	3.503213104	Up	7.09E-09	1.42E-08	AAEL001591-PA OS=Aedes aegypti GN=AAEL001591 PE=4 SV=1

CCG007502.1	573	23	223	1.72478865	13.60505361	2.979651148	Up	4.01E-11	8.67E-11	AAEL004909-PA OS=Aedes aegypti GN=AAEL004909 PE=4 SV=1
CCG022688.3	1572	23	223	0.62869205	4.95909397	2.979651148	Up	4.01E-11	8.67E-11	AAEL013154-PA OS=Aedes aegypti GN=AAEL013154 PE=4 SV=1
CCG004509.1	1661	66	223	1.70740667	4.693374907	1.458818984	Up	0	0	AAEL013993-PA OS=Aedes aegypti GN=AAEL013993 PE=4 SV=1
CCG021811.1	1691	72	223	1.82958066	4.610109829	1.333288102	Up	3.34E-13	9.11E-13	AAEL004456-PA OS=Aedes aegypti GN=AAEL004456 PE=4 SV=1
CCG001677.1	567	0	224	0.001	13.81067724	13.75349645	Up	4.44E-16	1.66E-15	5-hydroxyisourate hydrolase OS=Culex quinquefasciatus GN=CpipJ_CPIJ014057 PE=3 SV=1
CCG013598.1	1878	6	224	0.1372835	4.169677315	4.924705625	Up	8.38E-06	1.48E-05	AAEL004021-PA OS=Aedes aegypti GN=AAEL004021 PE=3 SV=1
CCG025997.1	1200	11	224	0.39388923	6.525544997	4.050236507	Up	2.67E-07	5.03E-07	AAEL013821-PA OS=Aedes aegypti GN=AAEL013821 PE=4 SV=1
CCG004958.2	954	16	224	0.72066641	8.208232701	3.509668126	Up	7.09E-09	1.42E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001384 PE=4 SV=1
CCG023427.1	1407	23	224	0.70241926	5.5654968	2.98610617	Up	4.01E-11	8.66E-11	AAEL000980-PA OS=Aedes aegypti GN=AAEL000980 PE=4 SV=1
CCG009730.1	582	54	224	3.98688259	13.45473195	1.754780624	Up	0	0	AAEL000050-PA OS=Aedes aegypti GN=AAEL000050 PE=4 SV=1
CCG004768.1	1392	63	224	1.94475092	5.625469825	1.532388202	Up	0	0	AGAP001675-PA OS=Anopheles gambiae GN=AgaP_AGAP001675 PE=4 SV=2
CCG015747.1	2161	81	224	1.61061939	3.623625172	1.169818123	Up	4.96E-11	1.07E-10	AAEL005676-PA OS=Aedes aegypti GN=AAEL005676 PE=4 SV=1
CCG004319.1	630	366	224	24.9633696	12.42960952	-1.006031712	Down	5.82E-17	2.20E-16	AAEL006702-PA OS=Aedes aegypti GN=AAEL006702 PE=4 SV=1
CCG001423.1	1225	390	224	13.6801604	6.39237061	-1.097662188	Down	1.66E-20	6.52E-20	Pyrroline-5-carboxylate reductase OS=Aedes aegypti GN=P5CR-1 PE=2 SV=1
CCG012954.1	888	624	224	30.1949486	8.818304051	-1.775734093	Down	1.60E-64	7.80E-64	AAEL006568-PA OS=Aedes aegypti GN=AAEL006568 PE=3 SV=1

CCG016175.1	1476	658	224	19.1558844	5.305321136	-1.852275648	Down	5.95E-72	2.96E-71	AAEL002672-PA OS=Aedes aegypti GN=AAEL002672 PE=4 SV=1
CCG025969.1	6015	677	224	4.83632757	1.301854364	-1.893343898	Down	3.46E-76	1.74E-75	Ubiquitin specific proteinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005165 PE=4 SV=1
CCG010652.1	1941	783	224	17.3340042	4.034340029	-2.103200371	Down	8.49E-101	4.50E-100	AAEL000223-PA OS=Aedes aegypti GN=AAEL000223 PE=4 SV=1
CCG000080.1	2019	1949	224	41.4799469	3.878481425	-3.418850249	Down	0	0	AAEL001813-PA OS=Aedes aegypti GN=AAEL001813 PE=3 SV=1
CCG022141.1	546	2013	224	158.421384	14.34185714	-3.465463331	Down	0	0	AAEL009872-PA OS=Aedes aegypti GN=AAEL009872 PE=4 SV=1
CCG009015.1	1050	17	225	0.69570047	7.491059308	3.428631554	Up	3.39E-09	6.89E-09	AAEL014658-PA OS=Aedes aegypti GN=AAEL014658 PE=3 SV=1
CCG019540.1	765	24	225	1.34807011	10.28184611	2.931131894	Up	1.93E-11	4.23E-11	AAEL008716-PA OS=Aedes aegypti GN=AAEL008716 PE=4 SV=1
CCG000210.1	625	48	225	3.30007562	12.58497964	1.931131894	Up	5.35E-14	1.79E-13	AAEL013608-PA OS=Aedes aegypti GN=AAEL013608 PE=4 SV=1
CCG021981.1	1359	61	225	1.92873717	5.787794168	1.585357057	Up	1.75E-14	6.17E-14	AAEL002493-PA OS=Aedes aegypti GN=AAEL002493 PE=3 SV=1
CCG003428.1	839	80	225	4.09723334	9.374984832	1.1941663	Up	2.15E-11	4.70E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_04538 PE=4 SV=1
CCG015766.1	2601	475	225	7.84722181	3.024072385	-1.375689308	Down	1.58E-34	6.84E-34	AAEL001707-PA OS=Aedes aegypti GN=AAEL001707 PE=4 SV=1
CCG013062.1	771	1805	225	100.597109	10.20183174	-3.301688727	Down	0	0	AAEL008159-PA OS=Aedes aegypti GN=AAEL008159 PE=4 SV=1
CCG017087.1	3615	3682	225	43.7661308	2.175826355	-4.330179516	Down	0	0	AAEL001766-PA OS=Aedes aegypti GN=AAEL001766 PE=4 SV=1
CCG022517.1	757	8	226	0.45410552	10.43668501	4.522492166	Up	2.19E-06	3.97E-06	AAEL000049-PA OS=Aedes aegypti GN=AAEL000049 PE=4 SV=1
CCG021982.1	996	29	226	1.25112681	7.93229975	2.664511171	Up	6.66E-13	1.69E-12	AAEL002493-PA OS=Aedes aegypti GN=AAEL002493 PE=3 SV=1
CCG024928.1	480	35	226	3.13320982	16.45952198	2.393209149	Up	6.39E-14	2.11E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_16009 PE=4 SV=1

CCG017637.1	2721	41	226	0.64746752	2.903554043	2.164940162	Up	1.02E-13	3.20E-13	AAEL011649-PA OS=Aedes aegypti GN=AAEL011649 PE=4 SV=1
CCG002583.1	657	66	226	4.31659434	12.02522154	1.478098047	Up	0	0	AAEL005414-PA OS=Aedes aegypti GN=AAEL005414 PE=4 SV=1
CCG023440.3	1665	69	226	1.78072774	4.745087418	1.413967709	Up	0	0	AAEL006419-PA OS=Aedes aegypti GN=AAEL006419 PE=4 SV=1
CCG009766.1	2589	70	226	1.16179275	3.051591561	1.393209149	Up	0	0	AAEL003070-PA OS=Aedes aegypti GN=AAEL003070 PE=4 SV=1
CCG015658.1	1857	1	227	0.02313933	4.273305777	7.528861691	Up	0.000587	0.00091615	AAEL013916-PA OS=Aedes aegypti GN=AAEL013916 PE=4 SV=1
CCG012891.1	1481	57	227	1.65379802	5.358223381	1.695971677	Up	0	0	AAEL009511-PA OS=Aedes aegypti GN=AAEL009511 PE=4 SV=1
CCG019590.1	1005	498	227	21.2924655	7.896048584	-1.431140241	Down	2.91E-38	1.29E-37	AAEL012025-PA OS=Aedes aegypti GN=AAEL012025 PE=4 SV=1
CCG006134.1	1008	0	228	0.001	7.90722927	12.94895654	Up	4.44E-16	1.65E-15	Mediator complex subunit 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ013633 PE=4 SV=1
CCG025309.1	1152	43	228	1.60390502	6.918825611	2.108938463	Up	8.46E-14	2.70E-13	AAEL006666-PA OS=Aedes aegypti GN=AAEL006666 PE=4 SV=1
CCG014093.1	774	51	228	2.8313391	10.29778696	1.862777876	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001626 PE=4 SV=1
CCG014548.1	2346	72	228	1.31876423	3.397479584	1.365278217	Up	2.35E-14	8.21E-14	AAEL006209-PA OS=Aedes aegypti GN=AAEL006209 PE=4 SV=1
CCG006484.1	999	81	228	3.48403254	7.97846557	1.195353215	Up	1.53E-11	3.39E-11	AAEL008794-PA OS=Aedes aegypti GN=AAEL008794 PE=4 SV=1
CCG006571.1	1137	85	228	3.21233724	7.010102994	1.125812282	Up	1.26E-10	2.69E-10	AAEL005732-PA OS=Aedes aegypti GN=AAEL005732 PE=3 SV=1
CCG024383.1	531	376	228	30.4267801	15.01033353	-1.019385634	Down	9.84E-18	3.75E-17	AAEL004496-PA OS=Aedes aegypti GN=AAEL004496 PE=4 SV=1
CCG009419.1	1851	702	228	16.2964634	4.306043816	-1.920124002	Down	1.99E-80	1.01E-79	AAEL006434-PA OS=Aedes aegypti GN=AAEL006434 PE=3 SV=1
CCG017389.1	2838	1170	228	17.7147955	2.808487352	-2.657089596	Down	2.11E-198	1.26E-197	UDP-glucuronosyl transferase OS=Aedes aegypti PE=2 SV=1

CCG007023.1	2072	1742	228	36.1260993	3.846760185	-3.231325691	Down	0	0	AAEL007495-PA OS=Aedes aegypti GN=AAEL007495 PE=3 SV=1
CCG008915.1	1455	10	229	0.29532464	5.502024317	4.219588897	Up	5.43E-07	1.01E-06	AAEL005325-PA OS=Aedes aegypti GN=AAEL005325 PE=4 SV=1
CCG004863.1	1947	27	229	0.5958823	4.111682271	2.78662949	Up	2.28E-12	5.41E-12	AAEL001984-PA OS=Aedes aegypti GN=AAEL001984 PE=4 SV=1
CCG019686.1	943	40	229	1.82268227	8.489337626	2.219588897	Up	6.86E-14	2.24E-13	AAEL005592-PA OS=Aedes aegypti GN=AAEL005592 PE=4 SV=1
CCG008715.1	1287	41	229	1.3688882	6.220237281	2.183964987	Up	1.02E-13	3.20E-13	AAEL008599-PA (Fragment) OS=Aedes aegypti GN=AAEL008599 PE=3 SV=1
CCG003476.1	2278	53	229	0.99973483	3.514242924	1.813596537	Up	0	0	AAEL012229-PA OS=Aedes aegypti GN=AAEL012229 PE=4 SV=1
CCG022577.1	1296	62	229	2.05565088	6.177041189	1.587320682	Up	0	0	AAEL005746-PA OS=Aedes aegypti GN=AAEL005746 PE=4 SV=1
CCG000878.1	1536	68	229	1.90230596	5.211878503	1.454054151	Up	0	0	SYB1-POZ domain-containing protein 3 OS=Culex quinquefasciatus GN=Cuq3.12-PA OS=Aedes aegypti GN=AAEL009412 PE=4 SV=1
CCG001176.1	2604	92	229	1.51813194	3.074287781	1.017955036	Up	2.83E-09	5.76E-09	AAEL009412-PA OS=Aedes aegypti GN=AAEL009412 PE=4 SV=1
CCG016487.1	3400	440	229	5.56078918	2.354542759	-1.239842722	Down	1.53E-27	6.33E-27	AAEL008242-PA OS=Aedes aegypti GN=AAEL008242 PE=4 SV=1
CCG005473.1	3422	527	229	6.61748981	2.33940543	-1.50014216	Down	3.07E-43	1.39E-42	AAEL004509-PA OS=Aedes aegypti GN=AAEL004509 PE=4 SV=1
CCG025116.1	876	36	230	1.7658795	9.178542988	2.377878253	Up	6.13E-14	2.03E-13	AAEL003484-PA OS=Aedes aegypti GN=AAEL003484 PE=4 SV=1
CCG024610.1	393	43	230	4.70152313	20.45904239	2.1215385	Up	8.46E-14	2.70E-13	AAEL013061-PA OS=Aedes aegypti GN=AAEL013061 PE=4 SV=1
CCG011650.1	1203	45	230	1.60734668	6.683627313	2.055950158	Up	0	0	AAEL004843-PA OS=Aedes aegypti GN=AAEL004843 PE=3 SV=1
CCG011865.1	1008	65	230	2.77086582	7.97659093	1.525435442	Up	0	0	AAEL011367-PA OS=Aedes aegypti GN=AAEL011367 PE=4 SV=1
CCG010516.1	1434	73	230	2.18744116	5.606976051	1.357978696	Up	0	0	AAEL009786-PA OS=Aedes aegypti GN=AAEL009786 PE=4 SV=1

CCG016958.1	1251	75	230	2.57612318	6.427181181	1.318984564	Up	1.19E-13	3.68E-13	AAEL004497-PA OS=Aedes aegypti GN=AAEL004497 PE=4 SV=1
CCG021350.1	706	3800	230	231.281858	11.38867374	-4.343980448	Down	0	0	AAEL002889-PA OS=Aedes aegypti GN=AAEL002889 PE=4 SV=1
CCG005677.1	408	0	231	0.001	19.79255376	14.27267015	Up	4.44E-16	1.63E-15	AAEL013744-PA OS=Aedes aegypti GN=AAEL013744 PE=2 SV=1
CCG012213.2	8548	15	231	0.07540314	0.94470776	3.64717165	Up	1.48E-08	2.93E-08	Polyprotein OS=Culex sinensis GN=CLF_101279 PE=4 SV=1
CCG025980.1	4288	30	231	0.30062781	1.88324672	2.64717165	Up	3.71E-13	9.94E-13	AAEL001966-PA OS=Aedes aegypti GN=AAEL001966 PE=4 SV=1
CCG027208.2	1080	39	231	1.55168486	7.477186976	2.268660026	Up	7.33E-14	2.38E-13	AAEL009134-PA (Fragment) OS=Aedes aegypti GN=AAEL009134 PE=4 SV=1
CCG027067.1	366	49	231	5.75277868	22.06383042	1.939352401	Up	0	0	Single-stranded DNA-binding protein OS=Aedes aegypti GN=AAEL000147 PE=3 SV=1
CCG028124.1	606	84	231	5.95620084	13.32567976	1.161744822	Up	3.17E-11	6.89E-11	AAEL010423-PA OS=Aedes aegypti GN=AAEL010423 PE=4 SV=1
CCG005101.1	1860	444	231	10.2572915	4.341592438	-1.240353621	Down	8.55E-28	3.55E-27	AAEL006905-PA OS=Aedes aegypti GN=AAEL006905 PE=4 SV=1
CCG006412.1	2932	559	231	8.19238801	2.754216212	-1.572642228	Down	5.87E-49	2.72E-48	AAEL010676-PA (Fragment) OS=Aedes aegypti GN=AAEL010676 PE=4 SV=1
CCG027941.1	1713	626	231	15.7028919	4.714163418	-1.735956602	Down	9.69E-63	4.71E-62	AAEL002880-PA OS=Aedes aegypti GN=AAEL002880 PE=3 SV=1
CCG025347.1	354	907	231	110.094772	22.81175688	-2.270896495	Down	3.01E-128	1.67E-127	AAEL002526-PA OS=Aedes aegypti GN=AAEL002526 PE=4 SV=1
CCG022765.1	1855	17	232	0.39379272	4.372140276	3.472831358	Up	3.39E-09	6.89E-09	AAEL010136-PA OS=Aedes aegypti GN=AAEL010136 PE=4 SV=1
CCG023408.1	828	20	232	1.03791629	9.795072719	3.238366104	Up	3.69E-10	7.71E-10	AAEL010949-PA OS=Aedes aegypti GN=AAEL010949 PE=4 SV=1
CCG011479.1	2420	27	232	0.47941439	3.351371988	2.805406697	Up	2.28E-12	5.41E-12	AAEL000918-PA (Fragment) OS=Aedes aegypti GN=AAEL000918 PE=4 SV=2
CCG008611.1	575	43	232	3.21338885	14.10490472	2.134029444	Up	8.46E-14	2.71E-13	-

CCG006115.1	1376	69	232	2.15473233	5.89412806	1.451769742	Up	0	0	AAEL002285-PA (Fragment) OS=Aedes aegypti GN=AAEL002285 PE=4 SV=1
CCG027355.1	5001	418	232	3.5915515	1.621739694	-1.147064933	Down	2.30E-23	9.21E-23	AAEL009135-PA (Fragment) OS=Aedes aegypti GN=AAEL009135 PE=4 SV=1
CCG012374.1	1008	574	232	24.4688767	8.04595259	-1.604612728	Down	1.12E-51	5.24E-51	AAEL000629-PA OS=Aedes aegypti GN=AAEL000629 PE=3 SV=1
CCG020204.1	503	2283	232	195.02963	16.12389704	-3.596420945	Down	0	0	-
CCG026165.1	375	0	233	0.001	21.72074263	14.40678581	Up	4.44E-16	1.64E-15	AAEL013544-PA OS=Aedes aegypti GN=AAEL013544 PE=4 SV=1
CCG003214.1	594	0	233	0.001	13.71259005	13.74321347	Up	4.44E-16	1.63E-15	AAEL000692-PA OS=Aedes aegypti GN=AAEL006892 PE=4 SV=1
CCG026680.1	309	7	233	0.97342441	26.36012456	4.759144426	Up	4.31E-06	7.71E-06	AAEL003968-PA OS=Aedes aegypti GN=AAEL003968 PE=4 SV=1
CCG021948.1	1554	13	233	0.35946367	5.241491948	3.86605963	Up	6.34E-08	1.23E-07	AAEL002785-PA OS=Aedes aegypti GN=AAEL002785 PE=4 SV=1
CCG013854.1	1468	15	233	0.43906405	5.548554828	3.659608753	Up	1.48E-08	2.92E-08	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001406 PE=4 SV=1
CCG014823.1	438	44	233	4.31659434	18.59652623	2.10706773	Up	3.91E-14	1.34E-13	AAEL010249-PA OS=Aedes aegypti GN=AAEL010249 PE=4 SV=1
CCG006833.1	456	60	233	5.65391245	17.86245282	1.659608753	Up	0	0	AAEL015541-PA (Fragment) OS=Aedes aegypti GN=AAEL015541 PE=4 SV=1
CCG007693.1	867	66	233	3.27105246	9.394784877	1.522105229	Up	0	0	AAEL012523-PA OS=Aedes aegypti GN=AAEL012523 PE=4 SV=1
CCG016298.1	1333	73	233	2.35318127	6.110486488	1.37667479	Up	0	0	AAEL005958-PA OS=Aedes aegypti GN=AAEL005958 PE=4 SV=1
CCG008797.1	2846	84	233	1.2682564	2.862009307	1.174181926	Up	1.77E-11	3.90E-11	-
CCG008454.1	1341	93	233	2.98000397	6.074033175	1.027340537	Up	1.56E-09	3.20E-09	AAEL015014-PA OS=Aedes aegypti GN=AAEL015014 PE=4 SV=1
CCG001528.1	747	1667	233	95.8909606	10.90398727	-3.136539041	Down	0	0	AAEL000629-PA OS=Aedes aegypti GN=AAEL006297 PE=4 SV=1

CCG005060.1	1050	22	234	0.90031825	7.790701681	3.113246305	Up	8.38E-11	1.80E-10	Fibrinogen and fibronectin OS=Culex quinquefasciatus GN=CpipJ_CPIJ009584 PE=4 SV=1
CCG007884.1	852	25	234	1.26084902	9.60121686	2.928821734	Up	9.32E-12	2.08E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006742 PE=4 SV=1
CCG021845.1	858	33	234	1.6526821	9.534075483	2.528283804	Up	1.31E-13	4.01E-13	AAEL012389-PA OS=Aedes aegypti GN=AAEL012389 PE=4 SV=1
CCG001508.2	2700	44	234	0.70024753	3.02971732	2.113246305	Up	3.91E-14	1.35E-13	AAEL003172-PA OS=Aedes aegypti GN=AAEL003172 PE=4 SV=1
CCG005642.2	1890	941	234	21.3939261	4.3281676	-2.305372989	Down	1.83E-135	1.02E-134	Uncharacterized protein OS=Anopheles darlingi GN=AND_22059 PE=4 SV=1
CCG006186.1	819	20	235	1.04932197	10.03076318	3.256902055	Up	3.69E-10	7.72E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012958 PE=4 SV=1
CCG025756.1	560	26	235	1.99502339	14.66999115	2.878390432	Up	4.60E-12	1.05E-11	-
CCG014001.1	249	41	235	7.07533783	32.99275117	2.221278146	Up	1.02E-13	3.20E-13	AAEL003749-PA OS=Aedes aegypti GN=AAEL003749 PE=4 SV=1
CCG017039.1	1425	91	235	2.74403218	5.765049152	1.07103551	Up	3.53E-10	7.41E-10	AAEL006300-PA OS=Aedes aegypti GN=AAEL006300 PE=4 SV=1
CCG023285.1	1374	468	235	14.6359795	5.979035692	-1.291534569	Down	5.08E-31	2.16E-30	AAEL006982-PA OS=Aedes aegypti GN=AAEL006982 PE=3 SV=1
CCG010990.1	2712	678	235	10.7424337	3.029201711	-1.826311313	Down	1.11E-72	5.52E-72	Neurexin-4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ011417 PE=4 SV=1
CCG001499.1	4503	783	235	7.47175265	1.824382643	-2.034038347	Down	1.51E-96	7.96E-96	AAEL014592-PA OS=Aedes aegypti GN=AAEL014592 PE=4 SV=1
CCG001431.2	14786	875	235	2.54284579	0.55560632	-2.194309056	Down	1.26E-118	6.87E-118	AGAP005010-PA OS=Anopheles gambiae GN=AGAP005010 PE=3 SV=4
CCG020885.1	1995	5920	235	127.509187	4.117892251	-4.95255131	Down	0	0	AAEL012082-PA OS=Aedes aegypti GN=AAEL012082 PE=4 SV=1

CCG006187.1	1935	14	236	0.31089214	4.263645126	3.777601331	Up	3.07E-08	6.01E-08	AAEL014096-PA (Fragment) OS=Aedes aegypti GN=AAEL014096 PE=4 SV=1
CCG020734.1	1373	23	236	0.71981347	6.008851652	3.061394297	Up	4.01E-11	8.68E-11	AAEL002159-PA OS=Aedes aegypti GN=AAEL002159 PE=4 SV=1
CCG018671.1	1647	30	236	0.78269098	5.009200558	2.678065658	Up	3.71E-13	9.95E-13	AAEL009725-PA OS=Aedes aegypti GN=AAEL009725 PE=4 SV=1
CCG016542.1	660	82	236	5.338664	12.5002323	1.227404249	Up	2.40E-12	5.67E-12	AAEL007525-PA OS=Aedes aegypti GN=AAEL007525 PE=3 SV=1
CCG003585.1	759	605	236	34.2512377	10.86976722	-1.655835079	Down	7.77E-57	3.71E-56	AAEL011004-PA OS=Aedes aegypti GN=AAEL011004 PE=4 SV=1
CCG011492.1	924	2249	236	104.58759	8.928737357	-3.550111693	Down	0	0	AAEL006065-PA OS=Aedes aegypti GN=AAEL006065 PE=3 SV=1
CCG002191.1	261	2	237	0.32927	31.74372259	6.591056453	Up	9.36E-05	0.0001537	AAEL012504-PA OS=Aedes aegypti GN=AAEL012504 PE=4 SV=1
CCG014347.1	1812	33	237	0.78256139	4.572357392	2.546662333	Up	1.31E-13	4.01E-13	AAEL014289-PA OS=Aedes aegypti GN=AAEL014289 PE=4 SV=1
CCG008882.1	984	52	237	2.27075833	8.419828857	1.890616735	Up	0	0	Methionine aminopeptidase OS=Aedes aegypti GN=AAEL003769 PE=3 SV=1
CCG017226.1	1414	57	237	1.73216045	5.859343419	1.758166439	Up	0	0	AAEL007864-PA OS=Aedes aegypti GN=AAEL007864 PE=4 SV=1
CCG025843.1	933	68	237	3.13177058	8.880076736	1.503593611	Up	0	0	AAEL008911-PA OS=Aedes aegypti GN=AAEL008911 PE=4 SV=1
CCG005105.1	441	72	237	7.01546688	18.78710112	1.421131451	Up	0	0	AAEL010292-PA (Fragment) OS=Aedes aegypti GN=AAEL010292 PE=4 SV=1
CCG002799.1	1077	73	237	2.91252612	7.692768426	1.401231894	Up	0	0	Sennelureonine protein kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ011728 PE=4 SV=1
CCG004547.1	2295	78	237	1.46040928	3.610070412	1.305654234	Up	6.28E-14	2.08E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_09268 PE=4 SV=1

CCG000399.1	897	473	237	22.6585111	9.236467776	-1.294639921	Down	1.91E-31	8.11E-31	Neurongin OS=Culex quinquefasciatus GN=CpipJ_CPIJ010245 PE=4 SV=1
CCG012644.1	672	672	237	42.9697346	12.32903511	-1.80126097	Down	1.03E-70	5.09E-70	AAEL013326-PA OS=Aedes aegypti GN=AAEL013326 PE=4 SV=1
CCG013054.1	1129	5	238	0.19029998	7.369415298	5.275202872	Up	1.59E-05	2.76E-05	Tetrancopeptide repeat protein, tpr OS=Culex quinquefasciatus GN=CpipJ_CPIJ016841 PE=4 SV=1
CCG000514.1	2381	12	238	0.21656313	3.494359459	4.012168466	Up	1.30E-07	2.49E-07	AAEL007904-PA OS=Aedes aegypti GN=AAEL007904 PE=4 SV=1
CCG005590.1	2148	41	238	0.82018581	3.873403106	2.239578963	Up	1.02E-13	3.20E-13	AAEL010786-PA OS=Aedes aegypti GN=AAEL010786 PE=4 SV=1
CCG015039.1	992	48	238	2.07918071	8.387167209	2.012168466	Up	5.35E-14	1.79E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009013 PE=4 SV=1
CCG026214.1	2916	443	238	6.52798094	2.853247555	-1.194031921	Down	3.35E-26	1.37E-25	AAEL004819-PA OS=Aedes aegypti GN=AAEL004819 PE=4 SV=1
CCG000611.1	921	487	238	22.7212386	9.033734931	-1.330646995	Down	1.15E-33	4.92E-33	AGAP007599-PA OS=Anopheles gambiae GN=AGAP007599 PE=4 SV=4
CCG021411.1	19747	1110	238	2.41537476	0.421333361	-2.519212994	Down	3.11E-177	1.82E-176	AAEL013387-PA OS=Aedes aegypti GN=AAEL013387 PE=4 SV=1
CCG021961.1	1824	1111	238	26.172903	4.561441816	-2.520512134	Down	1.70E-177	9.91E-177	AAEL014138-PA OS=Aedes aegypti GN=SRPN16 PE=3 SV=2
CCG013043.1	1083	0	239	0.001	7.714707432	12.91339573	Up	4.44E-16	1.63E-15	AAEL011245-PA OS=Aedes aegypti GN=AAEL011245 PE=4 SV=1
CCG027224.1	819	9	239	0.47219489	10.20149957	4.43325501	Up	1.09E-06	2.01E-06	-
CCG014330.1	1590	43	239	1.16207458	5.254734685	2.176915257	Up	8.46E-14	2.70E-13	AAEL013892-PA OS=Aedes aegypti GN=AAEL013892 PE=4 SV=1
CCG006385.2	2572	65	239	1.08593808	3.248455734	1.580812199	Up	0	0	AAEL008539-PA OS=Aedes aegypti GN=AAEL008539 PE=4 SV=1
CCG011430.1	912	67	239	3.15676778	9.161215075	1.537090821	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_19886 PE=4 SV=1

CCG010236.1	644	75	239	5.00423928	12.97364619	1.374361321	Up	0	0	AAEL008067-PA OS=Aedes aegypti GN=AAEL008067 PE=4 SV=1
CCG008418.1	2955	414	239	6.02012526	2.827420693	-1.090306946	Down	1.88E-21	7.44E-21	B-cell CLL/lymphoma 11A OS=Culex quinquefasciatus GN=CpipJ_CPIJ011139 PE=4 SV=1
CCG005782.1	491	479	239	41.9195578	17.01635061	-1.300701834	Down	4.80E-32	2.05E-31	AAEL006951-PA OS=Aedes aegypti GN=AAEL006951 PE=4 SV=1
CCG008748.1	1410	574	239	17.4926437	5.925551878	-1.561726915	Down	9.93E-50	4.63E-49	Tubulin beta-1 chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ003635 PE=3 SV=1
CCG011363.1	4227	676	239	6.87190457	1.976585793	-1.797699424	Down	6.21E-71	3.08E-70	AAEL005886-PA OS=Aedes aegypti GN=AAEL005886 PE=4 SV=1
CCG023919.1	900	877	239	41.8716192	9.28336461	-2.173253021	Down	1.84E-117	1.00E-116	AAEL009750-PA OS=Aedes aegypti GN=AAEL009750 PE=4 SV=1
CCG015716.1	9328	1956	239	9.01037745	0.895693412	-3.330510643	Down	0	0	-
CCG018297.1	474	37	240	3.3541776	17.7003933	2.399750434	Up	5.17E-14	1.74E-13	AAEL013696-PA OS=Aedes aegypti GN=AAEL013696 PE=4 SV=1
CCG017264.1	993	51	240	2.2069048	8.449130338	1.936778457	Up	0	0	AAEL006856-PA OS=Aedes aegypti GN=AAEL006856 PE=4 SV=1
CCG007940.1	1119	61	240	2.34240734	7.497753731	1.678466462	Up	1.75E-14	6.17E-14	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007533 PE=4 SV=1
CCG001547.1	2492	63	240	1.08631352	3.366768228	1.631923876	Up	0	0	AAEL009415-PA OS=Aedes aegypti GN=AAEL009415 PE=4 SV=1
CCG020955.1	1533	64	240	1.79390934	5.472920043	1.609203799	Up	0	0	AAEL008174-PA OS=Aedes aegypti GN=AAEL008174 PE=4 SV=1
CCG016192.1	497	78	240	6.74374105	16.88126041	1.323801581	Up	0	0	AAEL010572-PA OS=Aedes aegypti GN=AAEL010572 PE=4 SV=1
CCG012156.1	1017	84	240	3.54912262	8.249740831	1.216886377	Up	2.21E-12	5.26E-12	AAEL004827-PA OS=Aedes aegypti GN=AAEL004827 PE=1 SV=1
CCG003049.1	2895	86	240	1.27647571	2.898095484	1.182939045	Up	6.51E-12	1.46E-11	AAEL011258-PA OS=Aedes aegypti GN=AAEL011258 PE=4 SV=1
CCG003000.1	495	96	240	8.33352429	16.94946753	1.024241299	Up	9.72E-10	2.01E-09	-
CCG015530.1	1671	401	240	10.3117077	5.020937418	-1.038254627	Down	2.43E-19	9.41E-19	AAEL012287-PA OS=Aedes aegypti GN=AAEL012287 PE=3 SV=1

CCG007702.1	3129	426	240	5.85014604	2.681363511	-1.125505821	Down	4.12E-23	1.65E-22	Growth factor receptor-bound protein OS=Culex quinquefasciatus AAEL010793-PA OS=Aedes aegypti GN=AAEL010793 PE=4 SV=1
CCG027705.1	1545	1064	240	29.592102	5.430411926	-2.446078636	Down	3.29E-164	1.90E-163	AAEL003861-PA OS=Aedes aegypti GN=AAEL003861 PE=4 SV=1
CCG020999.1	795	3	241	0.16214994	10.59741472	6.030240039	Up	5.28E-05	8.84E-05	AAEL003861-PA OS=Aedes aegypti GN=AAEL003861 PE=4 SV=1
CCG021077.1	500	20	241	1.71878938	16.8498894	3.293274445	Up	3.69E-10	7.72E-10	-
CCG016152.1	879	26	241	1.27100466	9.584692494	2.914762822	Up	4.60E-12	1.06E-11	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015175 PE=4 SV=1
CCG004572.1	1095	39	241	1.5304289	7.694013427	2.329800321	Up	7.33E-14	2.38E-13	AAEL013006-PA OS=Aedes aegypti GN=AAEL013006 PE=4 SV=1
CCG006766.1	1646	51	241	1.33138303	5.11843542	1.942777198	Up	0	0	AAEL001359-PA OS=Aedes aegypti GN=AAEL001359 PE=4 SV=1
CCG013578.1	2127	53	241	1.07070801	3.960951905	1.887282085	Up	0	0	AAEL007932-PA OS=Aedes aegypti GN=AAEL007932 PE=4 SV=1
CCG018380.1	2770	73	241	1.13241539	3.041496282	1.425377981	Up	0	0	AAEL001530-PA OS=Aedes aegypti GN=AAEL001530 PE=4 SV=1
CCG003122.1	1749	83	241	2.03915836	4.817006691	1.240163109	Up	1.07E-12	2.64E-12	Cholinesterase OS=Culex quinquefasciatus GN=CpipJ_CPIJ019485 PE=4 SV=1
CCG026061.1	495	92	241	7.98629411	17.02009031	1.091640584	Up	1.10E-10	2.35E-10	AAEL010516-PA OS=Aedes aegypti GN=AAEL010516 PE=4 SV=1
CCG016840.1	1860	459	241	10.6038216	4.529540162	-1.227147803	Down	3.18E-28	1.33E-27	AAEL001368-PA OS=Aedes aegypti GN=AAEL001368 PE=4 SV=1
CCG000963.1	2574	661	241	11.0345744	3.27309429	-1.753303921	Down	4.62E-67	2.27E-66	AAEL000515-PA OS=Aedes aegypti GN=AAEL000515 PE=4 SV=1
CCG008462.1	1411	882	241	26.8598908	5.970903403	-2.169432306	Down	7.36E-118	4.02E-117	AAEL001920-PA OS=Aedes aegypti GN=AAEL001920 PE=4 SV=1
CCG003083.1	1504	11	242	0.31427332	5.624935491	4.161744822	Up	2.67E-07	5.04E-07	AAEL013616-PA OS=Aedes aegypti GN=AAEL013616 PE=4 SV=1
CCG027628.1	1395	35	242	1.0780937	6.06444658	2.491893424	Up	6.39E-14	2.11E-13	AAEL007758-PA OS=Aedes aegypti GN=AAEL007758 PE=4 SV=1

CCG018222.1	894	49	242	2.35516442	9.462978724	2.006466597	Up	0	0	AGAP001238-PB OS=Anopheles gambiae GN=AgaP_AGAP001238 PE=4 SV=1
CCG016579.1	537	53	242	4.24096077	15.75400927	1.893255987	Up	0	0	AAEL010088-PA OS=Aedes aegypti GN=AAEL010088 PE=4 SV=1
CCG009715.1	942	73	242	3.32992636	8.980788725	1.431351882	Up	0	0	AAEL002362-PA OS=Aedes aegypti GN=AAEL002362 PE=4 SV=1
CCG025600.1	1092	475	242	18.6910476	7.7471639	-1.270607262	Down	1.05E-30	4.44E-30	Lipase OS=Aedes aegypti GN=AAEL015326 PE=3 SV=1
CCG013038.1	4662	516	242	4.75598092	1.814651004	-1.390050814	Down	6.18E-38	2.73E-37	AAEL011241-PA OS=Aedes aegypti GN=AAEL011241 PE=4 SV=1
CCG006218.1	2257	17	243	0.3236533	3.76378434	3.539662866	Up	3.39E-09	6.90E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003716 PE=4 SV=1
CCG023066.1	1065	17	243	0.68590187	7.976395545	3.539662866	Up	3.39E-09	6.88E-09	AAEL004349-PA OS=Aedes aegypti GN=AAEL004349 PE=4 SV=1
CCG006991.1	1518	33	243	0.93412467	5.596087784	2.582731588	Up	1.31E-13	4.01E-13	AAEL007426-PA OS=Aedes aegypti GN=AAEL007426 PE=4 SV=1
CCG018489.1	618	48	243	3.33745512	13.74573019	2.042163207	Up	5.35E-14	1.79E-13	-
CCG008755.1	1962	64	243	1.40166311	4.32969483	1.627125707	Up	0	0	Leucine-rich transmembrane protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003712 PE=2 SV=1
CCG020594.1	1374	68	243	2.12659531	6.182577333	1.539662866	Up	0	0	AAEL010501-PA OS=Aedes aegypti GN=AAEL010501 PE=4 SV=1
CCG006741.1	567	91	243	6.89637716	14.98211862	1.119331067	Up	3.82E-11	8.27E-11	AAEL003678-PA OS=Aedes aegypti GN=AAEL003678 PE=3 SV=1
CCG027525.1	812	95	243	5.02724728	10.46165179	1.057270099	Up	2.73E-10	5.74E-10	AAEL012051-PA OS=Aedes aegypti GN=AAEL012051 PE=4 SV=1
CCG000919.1	1005	459	243	19.6249833	8.452598264	-1.215224636	Down	8.17E-28	3.39E-27	AAEL007500-PA OS=Aedes aegypti GN=AAEL007500 PE=4 SV=1
CCG024845.1	627	3942	243	270.154217	13.54842306	-4.317586354	Down	0	0	AAEL001323-PA OS=Aedes aegypti GN=AAEL001323 PE=4 SV=1
CCG017358.1	801	4	244	0.21458045	10.64896321	5.633050541	Up	2.95E-05	5.03E-05	-
CCG008095.1	877	26	244	1.27390319	9.726134016	2.932610823	Up	4.60E-12	1.05E-11	-

CCG027021.1	389	39	244	4.30801967	21.92755664	2.347648323	Up	7.33E-14	2.37E-13	AAEL008585-PA (Fragment) OS=Aedes aegypti GN=AAEL008585 PE=3 SV=1
CCG024129.1	1092	50	244	1.96747869	7.811190048	1.989194352	Up	0	0	AAEL004512-PA OS=Aedes aegypti GN=AAEL004512 PE=4 SV=1
CCG027732.1	645	79	244	5.26295974	13.22452641	1.329269793	Up	0	0	AAEL000944-PA OS=Aedes aegypti GN=AAEL000944 PE=4 SV=1
CCG024857.1	4049	546	244	5.79438753	2.10664844	-1.4597066	Down	5.27E-43	2.38E-42	AAEL002338-PA OS=Aedes aegypti GN=AAEL002338 PE=4 SV=1
CCG024710.1	1515	2	245	0.05672572	5.653318686	6.638951143	Up	9.36E-05	0.00015365	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG023277.1	2358	17	245	0.30979028	3.632221293	3.551488302	Up	3.39E-09	6.89E-09	AAEL008509-PA OS=Aedes aegypti GN=AAEL008509 PE=3 SV=1
CCG025198.1	1605	26	245	0.69608293	5.336310162	2.938511425	Up	4.60E-12	1.05E-11	AAEL002349-PA OS=Aedes aegypti GN=AAEL002349 PE=4 SV=1
CCG009070.1	1113	45	245	1.73732081	7.695218157	2.147098046	Up	0	0	AAEL007627-PA OS=Aedes aegypti GN=AAEL007627 PE=4 SV=1
CCG009769.1	819	46	245	2.41344053	10.45760416	2.115389187	Up	3.40E-14	1.18E-13	AAEL006720-PA OS=Aedes aegypti GN=AAEL006720 PE=4 SV=1
CCG011414.1	399	52	245	5.60006566	21.46560854	1.938511425	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014272 PE=4 SV=1
CCG028577.1	977	83	245	3.65044828	8.766405127	1.263911711	Up	4.09E-13	1.09E-12	AAEL011383-PA OS=Aedes aegypti GN=AAEL011383 PE=4 SV=1
CCG001788.1	716	91	245	5.46123722	11.96198018	1.131156503	Up	2.16E-11	4.73E-11	AAEL002758-PA OS=Aedes aegypti GN=AAEL002758 PE=4 SV=1
CCG026364.1	679	99	245	6.26510122	12.61381121	1.009594523	Up	1.03E-09	2.12E-09	AAEL009538-PA OS=Aedes aegypti GN=AAEL009538 PE=4 SV=1
CCG007821.1	1311	658	245	21.5668081	6.533011296	-1.722992631	Down	3.55E-65	1.73E-64	AAEL001876-PA OS=Aedes aegypti GN=AAEL001876 PE=3 SV=1
CCG003613.1	1125	1376	245	52.5567598	7.61313583	-2.787313612	Down	1.32E-245	8.12E-245	AAEL010793-PA OS=Aedes aegypti GN=AAEL010793 PE=4 SV=1
CCG009387.1	708	12	246	0.72830059	12.1465199	4.059865208	Up	1.30E-07	2.49E-07	AAEL003456-PA OS=Aedes aegypti GN=AAEL003456 PE=4 SV=1
CCG012074.1	739	15	246	0.87218676	11.63699064	3.737937114	Up	1.48E-08	2.92E-08	-

CCG000823.1	606	24	246	1.70177167	14.19098364	3.059865208	Up	1.93E-11	4.24E-11	DNA replication complex GINS protein PSF2 OS=Aedes aegypti GN=AAEL006660 PE=3 SV=1
CCG025809.1	1948	75	246	1.6543789	4.414648915	1.416009019	Up	0	0	AAEL004609-PA OS=Aedes aegypti GN=AAEL004609 PE=4 SV=1
CCG018526.1	579	1652	246	122.601039	14.85273935	-3.045170262	Down	0	0	-
CCG006309.1	1122	3	247	0.11489234	7.695806027	6.065717935	Up	5.28E-05	8.85E-05	-
CCG013531.1	264	4	247	0.65105659	32.70717562	5.650680435	Up	2.95E-05	5.03E-05	AAEL010910-PA OS=Aedes aegypti GN=AAEL010910 PE=4 SV=1
CCG018168.1	582	33	247	2.43642825	14.83624461	2.606286316	Up	1.31E-13	4.01E-13	AAEL002427-PA OS=Aedes aegypti GN=AAEL002427 PE=4 SV=1
CCG016656.1	924	89	247	4.13885972	9.344907319	1.174947004	Up	4.28E-12	9.85E-12	AAEL012438-PA OS=Aedes aegypti GN=AAEL012438 PE=4 SV=1
CCG005234.1	1758	448	247	10.950194	4.911657772	-1.156674487	Down	3.04E-25	1.24E-24	AAEL003139-PA OS=Aedes aegypti GN=AAEL003139 PE=4 SV=1
CCG004441.1	1755	541	247	13.245941	4.920053768	-1.428804348	Down	2.55E-41	1.14E-40	AAEL010462-PA OS=Aedes aegypti GN=AAEL010462 PE=4 SV=1
CCG010077.1	1399	617	247	18.9509123	6.172047436	-1.618446244	Down	4.27E-56	2.03E-55	AAEL011334-PA OS=Aedes aegypti GN=AAEL011334 PE=4 SV=1
CCG027055.1	1548	1026	247	28.4799404	5.577967935	-2.35213458	Down	3.57E-151	2.03E-150	AAEL015058-PA OS=Aedes aegypti GN=AAEL015058 PE=4 SV=1
CCG017077.1	1596	16	248	0.43077428	5.432113183	3.656509514	Up	7.09E-09	1.42E-08	Importin subunit alpha OS=Aedes aegypti GN=AAEL012960 PE=3 SV=1
CCG007058.1	246	75	248	13.1005288	35.2424904	1.427690824	Up	0	0	AGAP006038-PA OS=Anopheles gambiae GN=AGAP006038 PE=4 SV=4
CCG013318.1	1288	86	248	2.86909719	6.73109677	1.23024476	Up	4.58E-13	1.20E-12	AAEL004970-PA OS=Aedes aegypti GN=AAEL004970 PE=4 SV=1
CCG006344.1	570	96	248	7.23700793	15.20991691	1.071547013	Up	1.13E-10	2.41E-10	AAEL000467-PA OS=Aedes aegypti GN=AAEL000467 PE=2 SV=1
CCG001082.1	3114	439	248	6.05771146	2.784088837	-1.121567615	Down	1.21E-23	4.88E-23	AAEL009547-PA OS=Aedes aegypti GN=AAEL009547 PE=3 SV=1
CCG012801.1	566	860	248	65.2897028	15.31740749	-2.091683335	Down	9.44E-110	5.07E-109	AAEL002797-PA OS=Aedes aegypti GN=AAEL002797 PE=4 SV=1

CCG023354.1	1295	1120	248	37.1630137	6.694712463	-2.472773503	Down	5.67E-175	3.31E-174	Carbonic anhydrase OS=Aedes aegypti PE=2 SV=1
CCG005577.1	660	28	249	1.82295844	13.18880442	2.854960214	Up	1.17E-12	2.86E-12	AAEL007882-PA OS=Aedes aegypti GN=AAEL007882 PE=4 SV=1
CCG026616.1	426	51	249	5.144264	20.43335896	1.989889794	Up	0	0	AAEL004774-PA OS=Aedes aegypti GN=AAEL004774 PE=4 SV=1
CCG025764.1	2442	66	249	1.16134418	3.564541735	1.617921017	Up	0	0	AAEL010494-PA OS=Aedes aegypti GN=AAEL010494 PE=4 SV=1
CCG015966.1	1161	73	249	2.70180071	7.497511556	1.472490577	Up	0	0	AAEL007613-PA OS=Aedes aegypti GN=TOLL1A PE=4 SV=1
CCG014012.1	663	87	249	5.63856246	13.12912657	1.21937164	Up	6.66E-13	1.69E-12	AAEL004448-PA OS=Aedes aegypti GN=AAEL004448 PE=4 SV=1
CCG000405.1	1955	90	249	1.97814635	4.452486402	1.17046204	Up	3.94E-12	9.10E-12	AAEL004551-PA OS=Aedes aegypti GN=AAEL004551 PE=4 SV=1
CCG008773.1	1337	505	249	16.2301541	6.510554163	-1.317824442	Down	2.36E-34	1.02E-33	-
CCG018753.1	1773	531	249	12.8691083	4.909538024	-1.390252915	Down	5.36E-39	2.38E-38	AGAP002366-PA OS=Anopheles gambiae GN=AgaP_AGAP002366 PE=3 SV=3
CCG021125.1	3432	751	249	9.40275953	2.536308542	-1.890353962	Down	3.88E-84	1.98E-83	AAEL009195-PA OS=Aedes aegypti GN=AAEL009195 PE=4 SV=1
CCG013200.1	1887	838	249	19.0824789	4.612936363	-2.048491298	Down	3.76E-104	2.00E-103	AAEL006143-PA OS=Aedes aegypti GN=AAEL006143 PE=4 SV=1
CCG018777.1	1083	1	250	0.03967658	8.069777648	7.668097488	Up	0.000587	0.00091672	Arginine N-methyltransferase 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003726 PE=4 SV=1
CCG024576.1	474	7	250	0.63457414	18.43790969	4.860742566	Up	4.31E-06	7.71E-06	-
CCG005300.1	1257	20	250	0.68368711	6.952720122	3.346169394	Up	3.69E-10	7.71E-10	AAEL012371-PA OS=Aedes aegypti GN=AAEL012371 PE=4 SV=1
CCG021989.1	923	42	250	1.95528587	9.46865568	2.275780066	Up	7.66E-14	2.47E-13	-
CCG006329.1	2187	46	250	0.90379872	3.996145036	2.144535532	Up	3.40E-14	1.18E-13	AAEL007982-PA OS=Aedes aegypti GN=AAEL007982 PE=4 SV=1
CCG005144.1	1941	56	250	1.23972444	4.50261164	1.860742566	Up	0	0	AAEL014096-PA (Fragment) OS=Aedes aegypti GN=AAEL014096 PE=4 SV=1
CCG017433.1	524	75	250	6.15024827	16.67856716	1.439278798	Up	0	0	-

CCG019647.1	1947	87	250	1.92006518	4.488736103	1.225153993	Up	4.60E-13	1.20E-12	AAEL005423-PA OS=Aedes aegypti GN=AAEL005423 PE=4 SV=1
CCG009185.1	1065	1168	250	47.125493	8.206168256	-2.52172707	Down	1.31E-186	7.75E-186	AAEL005255-PA OS=Aedes aegypti GN=AAEL003255 PE=4 SV=1
CCG003157.1	636	3	251	0.20268743	13.79642684	6.088894257	Up	5.28E-05	8.83E-05	-
CCG006244.2	909	14	251	0.66180009	9.652945511	3.866501836	Up	3.07E-08	6.01E-08	AAEL012339-PA OS=Aedes aegypti GN=AAEL012339 PE=4 SV=1
CCG013486.1	717	27	251	1.61810716	12.23783469	2.918969256	Up	2.28E-12	5.41E-12	AAEL013815-PA OS=Aedes aegypti GN=AAEL013815 PE=4 SV=1
CCG023378.1	867	28	251	1.38771923	10.12056225	2.866501836	Up	1.17E-12	2.86E-12	-
CCG013827.1	1318	41	251	1.33669129	6.65745635	2.316304753	Up	1.02E-13	3.20E-13	AAEL011872-PA OS=Aedes aegypti GN=AAEL011872 PE=4 SV=1
CCG022014.1	1432	51	251	1.53034669	6.127463317	2.001431416	Up	0	0	AAEL013384-PA OS=Aedes aegypti GN=AAEL013384 PE=4 SV=1
CCG028419.1	918	87	251	4.07229511	9.558308791	1.230913262	Up	3.06E-13	8.44E-13	AAEL006923-PA OS=Aedes aegypti GN=AAEL006923 PE=4 SV=1
CCG000365.1	1893	826	251	18.7496042	4.635249588	-2.016141214	Down	1.39E-100	7.39E-100	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004327 PE=4 SV=1
CCG019795.1	1317	922	251	30.0820769	6.662511367	-2.174766183	Down	1.68E-123	9.26E-123	-
CCG013740.1	987	10	252	0.43535699	8.925517474	4.357665032	Up	5.43E-07	1.01E-06	AAEL008941-PA OS=Aedes aegypti GN=AAEL008941 PE=4 SV=1
CCG008215.1	906	15	252	0.71141945	9.723494202	3.772702532	Up	1.48E-08	2.93E-08	AAEL013096-PA OS=Aedes aegypti GN=AAEL013096 PE=4 SV=1
CCG006631.1	1227	73	252	2.55647158	7.179694985	1.489768568	Up	0	0	Schiffum OS=Culex quinquefasciatus GN=CpipJ_CPIJ000181 PE=4 SV=1
CCG026744.1	816	76	252	4.00208313	10.79593841	1.431665614	Up	0	0	AAEL000090-PA OS=Aedes aegypti GN=AAEL000090 PE=4 SV=1
CCG012458.1	789	84	252	4.5747246	11.16538117	1.287275705	Up	3.86E-14	1.33E-13	AAEL001778-PA OS=Aedes aegypti GN=AAEL001778 PE=4 SV=1
CCG020730.1	1830	930	252	21.8370782	4.813926637	-2.181493779	Down	4.84E-125	2.67E-124	AAEL009014-PA (Fragment) OS=Aedes aegypti GN=AAEL009014 PE=4 SV=1

CCG010159.1	838	1060	252	54.3531249	10.51251282	-2.370255422	Down	1.43E-157	8.19E-157	AAEL008697-PA OS=Aedes aegypti GN=AAEL008697 PE=4 SV=1
CCG016810.1	1644	34	253	0.88866848	5.379832131	2.597843937	Up	8.82E-14	2.80E-13	AAEL007446-PA OS=Aedes aegypti GN=AAEL007446 PE=4 SV=1
CCG021998.1	2848	45	253	0.67894595	3.105492986	2.193453682	Up	0	0	AAEL005690-PA OS=Aedes aegypti GN=AAEL005690 PE=4 SV=1
CCG002834.1	2151	48	253	0.95887832	4.111782438	2.100344278	Up	5.35E-14	1.79E-13	AAEL005472-PA OS=Aedes aegypti GN=AAEL005472 PE=4 SV=1
CCG026778.1	1361	52	253	1.64175327	6.498489363	1.98486706	Up	0	0	AAEL014554-PA OS=Aedes aegypti GN=AAEL014554 PE=4 SV=1
CCG018803.1	372	59	253	6.81509232	23.77538716	1.802663729	Up	0	0	AAEL005382-PA OS=Aedes aegypti GN=AAEL005382 PE=4 SV=1
CCG003560.1	10005	66	253	0.28345852	0.884002401	1.640912659	Up	0	0	Pharmin OS=Culex quinquefasciatus GN=Cole_CPI001694 PE=4 SV=1
CCG011989.1	2754	69	253	1.07658376	3.211490205	1.576782322	Up	0	0	AAEL003227-PA OS=Aedes aegypti GN=AAEL003227 PE=4 SV=1
CCG013337.1	531	79	253	6.3928607	16.65620343	1.38152603	Up	0	0	AAEL003794-PA OS=Aedes aegypti GN=AAEL003794 PE=3 SV=1
CCG004655.1	768	82	253	4.58791437	11.51620316	1.327754774	Up	0	0	AAEL013206-PA OS=Aedes aegypti GN=AAEL013206 PE=4 SV=1
CCG004645.1	2277	93	253	1.7550221	3.884252975	1.146147967	Up	6.25E-12	1.41E-11	AAEL007970-PA OS=Aedes aegypti GN=AAEL007970 PE=4 SV=1
CCG027108.1	1844	797	253	18.5720599	4.796336238	-1.953129135	Down	3.79E-93	1.98E-92	Cytochrome P450 OS=Aedes albopictus GN=CYP6Y7 PE=2 SV=1
CCG015921.1	819	979	253	51.3643104	10.79907695	-2.249858271	Down	1.07E-136	6.00E-136	Pyrroline-5-carboxylate reductase OS=Aedes aegypti GN=AAEL007033 PE=3 SV=1
CCG024712.1	711	2801	253	169.280206	12.43944307	-3.76641949	Down	0	0	AAEL011070-PA OS=Aedes aegypti GN=CTLGA3 PE=2 SV=1
CCG017672.1	1695	4208	253	106.676486	5.217961076	-4.353612211	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_09568 PE=4 SV=1
CCG008754.1	1560	7	254	0.19281291	5.691924551	4.883642969	Up	4.31E-06	7.70E-06	AAEL003543-PA OS=Aedes aegypti GN=AAEL003543 PE=4 SV=1
CCG017563.1	654	23	254	1.51116804	13.57706774	3.167435935	Up	4.01E-11	8.67E-11	-

CCG002954.1	821	39	254	2.04119324	10.81534994	2.405595672	Up	7.33E-14	2.38E-13	-
CCG015904.1	930	76	254	3.51150519	9.547744409	1.443070377	Up	0	0	AAEL011748-PA OS=Aedes aegypti GN=AAEL011748 PE=2 SV=1
CCG020874.1	234	81	254	14.8741389	37.94616368	1.351147888	Up	4.91E-14	1.66E-13	AAEL000211-PA OS=Aedes aegypti GN=AAEL000211 PE=4 SV=1
CCG018473.1	1591	85	254	2.29568035	5.581019673	1.281606954	Up	1.73E-14	6.10E-14	Transmembrane protein 15 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002894 PE=4 SV=1
CCG010868.1	414	86	254	8.92608014	21.44783164	1.264733136	Up	0	0	AAEL012108-PA OS=Aedes aegypti GN=AAEL012108 PE=4 SV=1
CCG008819.1	1743	441	254	10.8718606	5.094321457	-1.093636955	Down	7.48E-23	2.99E-22	AAEL000523-PA OS=Aedes aegypti GN=AAEL000523 PE=4 SV=1
CCG000591.1	939	492	254	22.5144935	9.456232482	-1.251516615	Down	4.91E-31	2.08E-30	Acyl-coenzyme A oxidase OS=Aedes aegypti GN=AAEL008574 PE=3 SV=1
CCG025944.1	2080	673	254	13.9031882	4.268943414	-1.703464804	Down	1.42E-65	6.93E-65	-
CCG023738.1	9026	696	254	3.31342071	0.983758287	-1.751945605	Down	1.93E-70	9.55E-70	AAEL013413-PA OS=Aedes aegypti GN=AAEL013413 PE=4 SV=1
CCG027444.1	1662	790	254	20.4248438	5.342600662	-1.934710952	Down	3.42E-91	1.78E-90	AAEL001517-PA OS=Aedes aegypti GN=AAEL001517 PE=4 SV=1
CCG004157.1	354	0	255	0.001	25.18180954	14.62009434	Up	4.44E-16	1.63E-15	-
CCG023859.1	1342	19	255	0.60836435	6.642593574	3.448739127	Up	7.73E-10	1.60E-09	AAEL014400-PA OS=Aedes aegypti GN=AAEL014400 PE=4 SV=1
CCG014095.1	633	33	255	2.24012834	14.08271813	2.652272521	Up	1.31E-13	4.01E-13	AAEL004624-PA OS=Aedes aegypti GN=AAEL004624 PE=4 SV=1
CCG010764.1	1079	71	255	2.82748022	8.261687282	1.546919521	Up	0	0	AAEL005149-PA (Fragment) OS=Aedes aegypti GN=AAEL005149 PE=4 SV=1
CCG003767.1	1986	90	255	1.94726894	4.488600492	1.204813544	Up	5.59E-13	1.44E-12	AAEL002870-PA OS=Aedes aegypti GN=AAEL002870 PE=4 SV=1
CCG001180.1	1193	92	255	3.3136761	7.472221774	1.173104685	Up	2.07E-12	4.93E-12	AAEL013336-PA OS=Aedes aegypti GN=AAEL013336 PE=4 SV=1
CCG007964.1	1986	97	255	2.09872319	4.488600492	1.096753798	Up	2.69E-11	5.86E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_05226 PE=4 SV=1

CCG004480.1	216	101	255	20.0923296	41.27018786	1.038455158	Up	1.83E-10	3.88E-10	AAEL002121-PA OS=Aedes aegypti GN=AAEL002121 PE=4 SV=1
CCG024395.1	1581	562	255	15.274504	5.638431737	-1.43775968	Down	2.99E-43	1.35E-42	Subunit-dependent prospanate transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ001942 PE=4 SV=1
CCG006400.1	1647	644	255	16.8017663	5.412483653	-1.634250237	Down	2.79E-59	1.34E-58	AAEL000311-PA OS=Aedes aegypti GN=AAEL000311 PE=4 SV=1
CCG023089.2	1365	1077	255	33.9035928	6.530667089	-2.376135894	Down	1.52E-160	8.73E-160	AAEL010480-PA OS=Aedes aegypti GN=AAEL010480 PE=1 SV=1
CCG027186.1	2555	41	256	0.6895339	3.502668827	2.344761199	Up	1.02E-13	3.19E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003369 PE=4 SV=1
CCG027957.1	813	49	256	2.5898118	11.00777227	2.08760336	Up	0	0	AAEL007983-PA OS=Aedes aegypti GN=AAEL007983 PE=4 SV=1
CCG008285.1	635	56	256	3.78945691	14.09341552	1.894958282	Up	0	0	AAEL000477-PA OS=Aedes aegypti GN=AAEL000477 PE=4 SV=1
CCG013673.1	573	82	256	6.14924649	15.61835751	1.344761199	Up	0	0	AAEL004623-PA OS=Aedes aegypti GN=AAEL004623 PE=4 SV=1
CCG019063.1	2685	86	256	1.3763118	3.333079648	1.276048449	Up	0	0	AAEL001217-PA OS=Aedes aegypti GN=AAEL001217 PE=4 SV=1
CCG017972.1	2472	711	256	12.3590135	3.620274617	-1.771392546	Down	5.06E-73	2.53E-72	AAEL004090-PA (Fragment) OS=Aedes aegypti GN=AAEL004090 PE=4 SV=1
CCG003998.1	1131	33	257	1.25375883	7.94365794	2.663543634	Up	1.31E-13	4.00E-13	AAEL007607-PA OS=Aedes aegypti GN=AAEL007607 PE=4 SV=1
CCG026365.1	653	41	257	2.69794658	13.75846421	2.350385748	Up	1.02E-13	3.20E-13	Stam1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ013090 PE=4 SV=1
CCG001170.1	1108	51	257	1.9778488	8.108553367	2.035512411	Up	0	0	AAEL009399-PA OS=Aedes aegypti GN=AAEL009399 PE=4 SV=1
CCG027769.1	2074	84	257	1.74033641	4.331859754	1.31562033	Up	0	0	Modifier of mag4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001651 PE=4 SV=1
CCG016761.1	832	530	257	27.3725473	10.79841001	-1.341910796	Down	5.97E-37	2.62E-36	AAEL001184-PA OS=Aedes aegypti GN=AAEL001184 PE=4 SV=1

CCG001470.1	1425	51	258	1.53786419	6.329288005	2.041115117	Up	0	0	AAEL014744-PA OS=Aedes aegypti GN=AAEL014744 PE=4 SV=1
CCG001544.1	189	86	258	19.552366	47.72082226	1.287275705	Up	0	0	AAEL009092-PA OS=Aedes aegypti GN=AAEL009092 PE=3 SV=1
CCG009027.1	1146	91	258	3.41208189	7.870187964	1.205745819	Up	3.90E-13	1.04E-12	AAEL012767-PA OS=Aedes aegypti GN=AAEL012767 PE=4 SV=1
CCG018496.1	795	558	258	30.1598892	11.3449502	-1.410580853	Down	8.97E-42	4.03E-41	AAEL011081-PA OS=Aedes aegypti GN=AAEL011081 PE=4 SV=1
CCG020018.1	8084	889	258	4.72539511	1.115689684	-2.08249915	Down	9.49E-113	5.12E-112	AAEL008307-PA OS=Aedes aegypti GN=AAEL008307 PE=4 SV=1
CCG001011.2	1284	1594	258	53.3440475	7.024326641	-2.924895455	Down	3.73E-299	2.35E-298	AAEL001232-PA OS=Aedes aegypti GN=AAEL001232 PE=3 SV=1
CCG016330.1	2752	5351	258	83.5505269	3.277338447	-4.672052354	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_16398 PE=4 SV=1
CCG021340.1	1038	5802	258	240.18343	8.689051452	-4.788794121	Down	0	0	AAEL000213-PA OS=Aedes aegypti GN=AAEL000213 PE=4 SV=1
CCG007437.1	603	32	259	2.28031759	15.01524657	2.719121492	Up	1.77E-13	5.17E-13	AAEL011589-PA OS=Aedes aegypti GN=AAEL011589 PE=4 SV=1
CCG003535.1	1412	38	259	1.15640929	6.412318473	2.471193978	Up	4.26E-14	1.45E-13	AAEL002045-PA OS=Aedes aegypti GN=AAEL002045 PE=4 SV=1
CCG005349.1	565	57	259	4.33499977	16.02512156	1.886231477	Up	0	0	AAEL003965-PA OS=Aedes aegypti GN=AAEL003965 PE=4 SV=1
CCG026819.1	691	72	259	4.47730954	13.10302993	1.54919649	Up	0	0	AAEL008656-PA OS=Aedes aegypti GN=AAEL008656 PE=4 SV=1
CCG000914.1	1493	77	259	2.21612161	6.064429795	1.452334951	Up	0	0	AAEL013054-PA OS=Aedes aegypti GN=AAEL013054 PE=4 SV=1
CCG008155.1	614	104	259	7.27826124	14.74624379	1.018681773	Up	2.54E-10	5.35E-10	GK21198 OS=Drosophila willistoni GN=GK21198 PE=4 SV=1
CCG025334.1	3315	457	259	5.92373114	2.731280146	-1.116928864	Down	2.16E-24	8.73E-24	AAEL011038-PA OS=Aedes aegypti GN=AAEL011038 PE=3 SV=1
CCG017782.2	2964	587	259	8.5098631	3.054721216	-1.478095202	Down	5.98E-47	2.75E-46	AAEL006825-PA OS=Aedes aegypti GN=AAEL006825 PE=4 SV=1

CCG010853.1	2025	732	259	15.5327633	4.471206758	-1.796578347	Down	1.36E-76	6.83E-76	AAEL004479-PA (Fragment) OS=Aedes aegypti GN=AAEL004479 PE=4 SV=1
CCG010648.1	425	1	260	0.10110526	21.38623991	7.724681017	Up	0.000587	0.00091859	Histone H2B OS=Aedes aegypti GN=AAEL015674 PE=3 SV=1
CCG014288.1	1413	4	260	0.12164115	6.43252085	5.724681017	Up	2.95E-05	5.04E-05	AAEL014380-PA OS=Aedes aegypti GN=AAEL014380 PE=4 SV=1
CCG013606.1	657	9	260	0.5886265	13.83432566	4.554756015	Up	1.09E-06	2.01E-06	-
CCG002977.1	932	34	260	1.56756543	9.752308971	2.637218176	Up	8.82E-14	2.80E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016896 PE=4 SV=1
CCG020716.1	2117	43	260	0.87279102	4.293411413	2.298416262	Up	8.46E-14	2.70E-13	AAEL003176-PA OS=Aedes aegypti GN=AAEL003176 PE=4 SV=1
CCG013447.1	1111	46	260	1.77912493	8.18105487	2.201119061	Up	3.40E-14	1.18E-13	AAEL003916-PA OS=Aedes aegypti GN=AAEL003916 PE=4 SV=1
CCG016511.1	500	59	260	5.07042868	18.17830392	1.842037967	Up	0	0	AAEL012167-PA OS=Aedes aegypti GN=AAEL012167 PE=4 SV=1
CCG008475.1	1042	68	260	2.80416694	8.722794588	1.637218176	Up	0	0	AAEL005638-PA OS=Aedes aegypti GN=AAEL005638 PE=4 SV=1
CCG027035.1	1509	99	260	2.81908796	6.023294871	1.095324397	Up	1.80E-11	3.96E-11	AAEL006353-PA OS=Aedes aegypti GN=AAEL006353 PE=4 SV=1
CCG007155.1	633	102	260	6.92403307	14.35884986	1.052255675	Up	7.66E-11	1.64E-10	AGAP010064-PA OS=Anopheles gambiae GN=AGAP010064 PE=4 SV=1
CCG013255.1	4449	586	260	5.65975826	2.042965152	-1.470075838	Down	1.67E-46	7.67E-46	AAEL006955-PA OS=Aedes aegypti GN=AAEL006955 PE=4 SV=1
CCG016535.1	4242	1828	260	18.5168965	2.142657228	-3.111369338	Down	0	0	AGAP007349-PA (Fragment) OS=Anopheles gambiae GN=AGAP007349 PE=4 SV=4
CCG008846.1	1476	10	261	0.29112286	6.181646502	4.408291106	Up	5.43E-07	1.01E-06	AAEL000980-PA OS=Aedes aegypti GN=AAEL000980 PE=4 SV=1
CCG021759.1	1427	14	261	0.42156712	6.393910468	3.922864278	Up	3.07E-08	6.01E-08	Terminal deoxycytidyl transferase rev1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007351 PE=4 SV=1

CCG003085.1	786	24	261	1.31205297	11.60828274	3.1452567	Up	1.93E-11	4.24E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010852 PE=4 SV=1
CCG013143.1	1426	35	261	1.05465688	6.398394276	2.600936183	Up	6.39E-14	2.11E-13	AAEL004040-PA OS=Aedes aegypti GN=AAEL004646 PE=3 SV=1
CCG017140.1	2352	86	261	1.57117227	3.87929857	1.303954446	Up	0	0	AAEL005631-PA OS=Aedes aegypti GN=AAEL005631 PE=4 SV=1
CCG026827.1	480	94	261	8.41490636	19.00856299	1.175630349	Up	9.14E-13	2.28E-12	AAEL009911-PA OS=Aedes aegypti GN=AAEL009911 PE=4 SV=1
CCG017982.1	954	689	261	31.0336972	9.564056853	-1.698140972	Down	8.46E-67	4.15E-66	AAEL004080-PA OS=Aedes aegypti GN=AAEL004086 PE=4 SV=1
CCG011722.1	1302	2675	261	88.2826729	7.007765159	-3.655103976	Down	0	0	AAEL013637-PA OS=Aedes aegypti GN=AAEL013637 PE=4 SV=1
CCG027270.1	642	23	262	1.53941417	14.26646186	3.212174249	Up	4.01E-11	8.66E-11	AAEL004121-PA OS=Aedes aegypti GN=AAEL004121 PE=4 SV=1
CCG005561.1	722	35	262	2.08302038	12.68569046	2.606453188	Up	6.39E-14	2.11E-13	-
CCG004020.1	1836	59	262	1.3808357	4.988599409	1.853093156	Up	0	0	AAEL005684-PA OS=Aedes aegypti GN=AAEL005684 PE=3 SV=1
CCG021099.1	1896	97	262	2.19834613	4.830732339	1.135823363	Up	3.78E-12	8.77E-12	AAEL001982-PA OS=Aedes aegypti GN=AAEL001982 PE=4 SV=1
CCG020665.1	3384	653	262	8.29173661	2.70658053	-1.615202976	Down	4.25E-59	2.04E-58	AAEL007670-PA (Fragment) OS=Aedes aegypti GN=AAEL007670 PE=4 SV=1
CCG015127.2	2805	659	262	10.0952068	3.265265068	-1.62839845	Down	2.58E-60	1.25E-59	AAEL005397-PA OS=Aedes aegypti GN=AAEL005397 PE=4 SV=1
CCG028096.1	3840	662	262	7.40780321	2.385174092	-1.634951201	Down	6.32E-61	3.06E-60	AAEL005906-PA OS=Aedes aegypti GN=AAEL005908 PE=4 SV=1
CCG026950.1	858	30	263	1.50243827	10.71564894	2.834341598	Up	3.71E-13	9.96E-13	AAEL004315-PA OS=Aedes aegypti GN=AAEL004315 PE=4 SV=1
CCG019079.1	1194	34	263	1.22359378	7.700189942	2.653769352	Up	8.82E-14	2.80E-13	AGAP004623-PA OS=Anopheles gambiae GN=AgaP_AGAP004623 PE=4 SV=5
CCG004076.1	894	42	263	2.01871236	10.2841463	2.34891477	Up	7.66E-14	2.47E-13	AAEL015062-PA OS=Aedes aegypti GN=AAEL015062 PE=4 SV=1

CCG015984.1	276	57	263	8.87418432	33.31169127	1.908342179	Up	0	0	AAEL010119-PA OS=Aedes aegypti GN=AAEL010119 PE=4 SV=1
CCG007396.1	390	83	263	9.14484096	23.57442767	1.366192762	Up	1.33E-13	4.03E-13	AAEL004069-PA OS=Aedes aegypti GN=AAEL004069 PE=4 SV=1
CCG012093.1	303	437	263	61.9728516	30.34332274	-1.030257276	Down	1.09E-20	4.26E-20	AAEL010095-PA (Fragment) OS=Aedes aegypti GN=AAEL010095 PE=2 SV=1
CCG025382.1	2224	567	263	10.9549638	4.134004852	-1.405972732	Down	3.23E-42	1.46E-41	AAEL004544-PA OS=Aedes aegypti GN=AAEL004544 PE=4 SV=1
CCG004025.1	2220	1007	263	19.4912265	4.14145351	-2.234615775	Down	2.27E-139	1.27E-138	CD109 antigen OS=Culex quinquefasciatus GN=CpipJ_CPIJ020143 PE=4 SV=1
CCG007800.1	1439	3006	263	89.7616555	6.389177756	-3.812397101	Down	0	0	AAEL015304-PA OS=Aedes aegypti GN=AAEL015304 PE=4 SV=1
CCG025369.1	765	29	264	1.62891804	12.06403277	2.888726328	Up	6.66E-13	1.69E-12	AAEL006603-PA OS=Aedes aegypti GN=AAEL006603 PE=4 SV=1
CCG027016.1	384	36	264	4.02841262	24.03381528	2.576782322	Up	6.13E-14	2.03E-13	AAEL000755-PA OS=Aedes aegypti GN=AAEL000755 PE=4 SV=1
CCG020605.1	1578	52	264	1.41598618	5.848532996	2.046267605	Up	0	0	AAEL002151-PA OS=Aedes aegypti GN=AAEL002151 PE=4 SV=1
CCG002757.1	1044	65	264	2.67531873	8.840024011	1.72433951	Up	0	0	AAEL005725-PA OS=Aedes aegypti GN=AAEL005725 PE=4 SV=1
CCG021717.1	2826	75	264	1.14038574	3.265741355	1.517888633	Up	0	0	AAEL005033-PA OS=Aedes aegypti GN=AAEL005033 PE=4 SV=1
CCG019217.1	1824	80	264	1.88463748	5.059750585	1.424779228	Up	0	0	AAEL011781-PA OS=Aedes aegypti GN=AAEL011781 PE=4 SV=1
CCG007343.1	489	1816	264	159.576765	18.8731801	-3.079841164	Down	0	0	AAEL009066-PA OS=Aedes aegypti GN=AAEL009066 PE=4 SV=1
CCG012028.1	318	3580	264	483.747327	29.02196562	-4.059036549	Down	0	0	AAEL013515-PA OS=Aedes aegypti GN=AAEL013515 PE=4 SV=1
CCG024095.1	564	26	265	1.98087429	16.42543146	3.051722035	Up	4.60E-12	1.05E-11	AAEL013814-PA OS=Aedes aegypti GN=AAEL013814 PE=4 SV=1
CCG005332.2	477	34	265	3.06283224	19.42126487	2.664698912	Up	8.82E-14	2.80E-13	AAEL013273-PA OS=Aedes aegypti GN=AAEL013273 PE=4 SV=1
CCG025603.1	532	48	265	3.87696854	17.41342734	2.167199253	Up	5.35E-14	1.80E-13	-

CCG015499.1	1340	50	265	1.60334831	6.913390556	2.108305563	Up	0	0	AAEL013869-PA OS=Aedes aegypti GN=AAEL013869 PE=4 SV=1
CCG021161.1	1482	54	265	1.56569883	6.250973917	1.997274251	Up	0	0	AAEL007981-PA OS=Aedes aegypti GN=AAEL007981 PE=4 SV=1
CCG020773.1	3720	60	265	0.69306024	2.490307351	1.845271158	Up	0	0	-
CCG026228.2	12870	440	265	1.46905076	0.719809118	-1.02919796	Down	8.65E-21	3.40E-20	AGAP005010-PA OS=Anopheles gambiae GN=AGAP005010 PE=3 SV=4
CCG018323.1	1523	480	265	13.5426609	6.082694251	-1.154728842	Down	7.32E-27	3.02E-26	AAEL003674-PA OS=Aedes aegypti GN=AAEL003674 PE=4 SV=1
CCG002976.1	678	11	266	0.69714909	13.71519413	4.298164021	Up	2.67E-07	5.03E-07	-
CCG021204.1	1425	21	266	0.63323819	6.525544997	3.365278217	Up	1.76E-10	3.72E-10	AAEL011893-PA OS=Aedes aegypti GN=AAEL011893 PE=4 SV=1
CCG013359.3	1536	33	266	0.92317789	6.05397241	2.71320152	Up	1.31E-13	4.02E-13	AAEL006827-PA OS=Aedes aegypti GN=CYP12F8 PE=3 SV=1
CCG005258.1	495	52	266	4.51399232	18.78565984	2.057155921	Up	0	0	AAEL011767-PA OS=Aedes aegypti GN=AAEL011767 PE=4 SV=1
CCG025633.1	1082	53	266	2.10480216	8.594178948	2.029675185	Up	0	0	AAEL006816-PA OS=Aedes aegypti GN=AAEL006816 PE=4 SV=1
CCG025459.1	1321	931	266	30.2837418	7.039289645	-2.105041718	Down	1.25E-119	6.83E-119	-
CCG018834.1	2271	19	267	0.3595002	4.110021972	3.515081622	Up	7.73E-10	1.60E-09	AGAP007560-PA OS=Anopheles gambiae GN=AGAP007560 PE=4 SV=4
CCG001406.1	993	23	267	0.99527079	9.399657501	3.239447179	Up	4.01E-11	8.66E-11	AAEL003004-PA OS=Aedes aegypti GN=AAEL003004 PE=4 SV=1
CCG011499.1	1596	25	267	0.67308482	5.848283144	3.119152946	Up	9.32E-12	2.08E-11	Purative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005826 PE=4 SV=1
CCG009841.1	1632	59	267	1.55344016	5.719276898	1.880366086	Up	0	0	AAEL010492-PA OS=Aedes aegypti GN=AAEL010492 PE=4 SV=1
CCG014799.2	1932	82	267	1.8237672	4.831190424	1.405457131	Up	0	0	Malate dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ016732 PE=4 SV=1
CCG004314.1	519	82	267	6.78905248	17.9843158	1.405457131	Up	0	0	AAEL006566-PA OS=Aedes aegypti GN=AAEL006566 PE=4 SV=1

CCG021657.1	1269	84	267	2.84433231	7.355287548	1.370691713	Up	0	0	AAEL010726-PA OS=Aedes aegypti GN=AAEL010728 PE=4 SV=1
CCG014867.1	5108	92	267	0.77392631	1.827302251	1.239447179	Up	7.15E-14	2.33E-13	AAEL014406-PA OS=Aedes aegypti GN=AAEL014406 PE=4 SV=1
CCG011051.1	1530	793	267	22.2712415	6.100562025	-1.86816792	Down	2.42E-87	1.25E-86	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009754 PE=4 SV=1
CCG016178.1	3477	922	267	11.3943328	2.684457837	-2.085613805	Down	4.33E-117	2.36E-116	AAEL012475-PA (Fragment) OS=Aedes aegypti GN=AAEL012475 PE=4 SV=1
CCG018942.1	1497	1183	267	33.9567108	6.235043352	-2.445225223	Down	2.78E-182	1.63E-181	AAEL009124-PA OS=Aedes aegypti GN=CYP6N12 PE=3 SV=1
CCG008299.1	759	107	268	6.05765692	12.34363396	1.026935408	Up	9.36E-11	2.00E-10	AAEL000625-PA OS=Aedes aegypti GN=AAEL006625 PE=4 SV=1
CCG014485.1	4287	468	268	4.69088775	2.185401954	-1.101962325	Down	1.96E-24	7.93E-24	AAEL004214-PA (Fragment) OS=Aedes aegypti GN=AAEL004214 PE=4 SV=1
CCG003334.1	971	583	268	25.799542	9.648628399	-1.418949679	Down	6.01E-44	2.73E-43	Uncharacterized protein OS=Anopheles darlingi GN=AND_16013 PE=4 SV=1
CCG018090.1	663	1033	268	66.9498278	14.13094747	-2.244222145	Down	1.10E-143	6.23E-143	AAEL007946-PA OS=Aedes aegypti GN=GSTE6 PE=3 SV=2
CCG018139.1	2086	1286	268	26.49045	4.491283881	-2.560272533	Down	6.84E-209	4.10E-208	AAEL000378-PA OS=Aedes aegypti GN=AAEL000378 PE=4 SV=1
CCG020041.1	1521	6	269	0.16950586	6.182627516	5.188813066	Up	8.38E-06	1.47E-05	AAEL004948-PA OS=Aedes aegypti GN=AAEL004948 PE=3 SV=1
CCG011109.1	549	8	269	0.62615278	17.12891886	4.773775566	Up	2.19E-06	3.96E-06	-
CCG000076.1	564	24	269	1.82849935	16.6733625	3.188813066	Up	1.93E-11	4.24E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ019728 PE=4 SV=1
CCG019721.1	2514	27	269	0.4614888	3.740563426	3.018888064	Up	2.28E-12	5.40E-12	Leucine-rich transmembrane protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011619 PE=4 SV=1
CCG009438.1	1203	33	269	1.1787209	7.816938031	2.729381447	Up	1.31E-13	4.00E-13	AAEL004843-PA OS=Aedes aegypti GN=AAEL004843 PE=3 SV=1

CCG008759.1	1041	66	269	2.72430594	9.033406774	1.729381447	Up	0	0	AAEL011634-PA OS=Aedes aegypti GN=AAEL011634 PE=4 SV=1
CCG004263.1	1341	74	269	2.37118595	7.012510404	1.564322201	Up	0	0	AGAP003454-PA OS=Anopheles gambiae GN=AgaP_AGAP003454 PE=4 SV=5
CCG003230.1	1674	88	269	2.25886299	5.617548657	1.314343948	Up	3.22E-14	1.12E-13	AAEL007981-PA OS=Aedes aegypti GN=AAEL007981 PE=4 SV=1
CCG025126.1	1602	101	269	2.70907815	5.870022754	1.115564084	Up	3.89E-12	9.00E-12	AAEL011900-PA OS=Aedes aegypti GN=AAEL011900 PE=4 SV=1
CCG001126.1	1512	104	269	2.95559021	6.21942887	1.073335848	Up	1.73E-11	3.80E-11	AAEL005739-PA (Fragment) OS=Aedes aegypti GN=AAEL005739 PE=4 SV=1
CCG003882.1	4256	1492	269	15.0636382	2.209533941	-2.769256254	Down	2.86E-264	1.77E-263	Adams-20 OS=Culex quinquefasciatus GN=CpipJ_CPIJ010188 PE=4 SV=1
CCG019554.1	1572	4	270	0.10933775	6.004284178	5.779128801	Up	2.95E-05	5.03E-05	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG016015.1	3642	23	270	0.27136296	2.591635016	3.255566845	Up	4.01E-11	8.68E-11	AAEL006754-PA OS=Aedes aegypti GN=AAEL006754 PE=4 SV=1
CCG013977.1	2476	26	270	0.45121692	3.812089955	3.078689083	Up	4.60E-12	1.05E-11	AAEL010401-PA OS=Aedes aegypti GN=AAEL010401 PE=4 SV=1
CCG028196.1	967	44	270	1.95518958	9.760842532	2.319697182	Up	3.91E-14	1.34E-13	AAEL010842-PA OS=Aedes aegypti GN=AAEL010842 PE=4 SV=1
CCG019857.1	1999	46	270	0.9887983	4.721728228	2.255566845	Up	3.40E-14	1.18E-13	AAEL010405-PA OS=Aedes aegypti GN=AAEL010405 PE=4 SV=1
CCG011395.1	2192	56	270	1.09776694	4.305992121	1.971773879	Up	0	0	AAEL000322-PA OS=Aedes aegypti GN=AAEL000322 PE=4 SV=1
CCG028486.1	2796	56	270	0.86062416	3.375799259	1.971773879	Up	0	0	AAEL004211-PA OS=Aedes aegypti GN=AAEL004211 PE=4 SV=1
CCG023913.3	2460	480	270	8.38433846	3.836884036	-1.127761795	Down	6.46E-26	2.65E-25	AAEL000549-PA OS=Aedes aegypti GN=AAEL000549 PE=4 SV=1
CCG001793.1	363	0	271	0.001	26.09832784	14.67166975	Up	4.44E-16	1.64E-15	-
CCG003330.1	1435	48	271	1.43731516	6.601876659	2.199499744	Up	5.35E-14	1.79E-13	AAEL004838-PA OS=Aedes aegypti GN=AAEL004838 PE=4 SV=1

CCG019597.1	754	78	271	4.44514496	12.56457958	1.499060026	Up	0	0	AAEL007357-PA OS=Aedes aegypti GN=AAEL007357 PE=4 SV=1
CCG026754.1	1197	87	271	3.12311354	7.914530497	1.341518749	Up	0	0	Gastrin zinc finger protein XLCGF52.1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006243 PE=4 SV=1
CCG013441.1	417	93	271	9.58317822	22.71868826	1.245303434	Up	7.04E-14	2.29E-13	AAEL000702-PA OS=Aedes aegypti GN=AAEL000702 PE=4 SV=1
CCG020768.2	19485	570	271	1.25700532	0.486204414	-1.370355864	Down	7.37E-41	3.30E-40	AAEL005929-PA OS=Aedes aegypti GN=AAEL005929 PE=3 SV=1
CCG021498.1	234	0	272	0.001	40.63526189	15.31044457	Up	4.44E-16	1.65E-15	Putative secreted protein (Precursor) OS=Aedes aegypti GN=SGP1A2 PE=4 SV=1
CCG025613.1	947	22	272	0.99824093	10.04081445	3.330344426	Up	8.38E-11	1.80E-10	AAEL008088-PA OS=Aedes aegypti GN=AAEL008088 PE=4 SV=1
CCG012587.1	1329	34	272	1.09930096	7.154741371	2.702313204	Up	8.82E-14	2.80E-13	Zinc finger and BTB domain-containing protein 24 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017225 PE=4 SV=1
CCG017962.1	627	66	272	4.52312996	15.16531305	1.745381926	Up	0	0	AAEL001162-PA OS=Aedes aegypti GN=AAEL001162 PE=4 SV=1
CCG011391.2	920	70	272	3.26943633	10.33549052	1.660493028	Up	0	0	AAEL000660-PA OS=Aedes aegypti GN=AAEL000660 PE=4 SV=1
CCG027658.1	1641	84	272	2.19954766	5.794424913	1.397458622	Up	0	0	AAEL002321-PA OS=Aedes aegypti GN=AAEL002321 PE=3 SV=1
CCG011459.1	693	96	272	5.95251735	13.72099752	1.204813544	Up	0	0	AAEL011952-PA OS=Aedes aegypti GN=AAEL011952 PE=4 SV=1
CCG010570.1	3591	607	272	7.26333303	2.647911802	-1.455776661	Down	1.98E-47	9.15E-47	AAEL012395-PA OS=Aedes aegypti GN=AAEL012395 PE=3 SV=1
CCG026119.1	1341	1352	272	43.3222082	7.09071684	-2.611103391	Down	1.70E-224	1.03E-223	AAEL010480-PA OS=Aedes aegypti GN=AAEL010480 PE=1 SV=1
CCG016350.1	1317	3836	272	125.157101	7.219932636	-4.11561096	Down	0	0	AAEL009863-PA OS=Aedes aegypti GN=AAEL009863 PE=4 SV=1
CCG012317.1	1203	35	273	1.25015853	7.933175028	2.665787328	Up	6.39E-14	2.11E-13	AAEL013688-PA OS=Aedes aegypti GN=AAEL013688 PE=4 SV=1

CCG024785.1	956	37	273	1.66305458	9.982855187	2.585616979	Up	5.17E-14	1.74E-13	AAEL008897-PA OS=Aedes aegypti GN=AAEL008897 PE=4 SV=1
CCG025437.1	1581	39	273	1.05997448	6.036438684	2.509668126	Up	7.33E-14	2.37E-13	AAEL008951-PA OS=Aedes aegypti GN=AAEL008951 PE=4 SV=1
CCG019259.1	1034	64	273	2.65963541	9.229796479	1.795070345	Up	0	0	AAEL014259-PA OS=Aedes aegypti GN=AAEL014259 PE=4 SV=1
CCG002866.6	1767	80	273	1.94543224	5.401024085	1.47314225	Up	0	0	Broad complex isoform Z1 OS=Aedes aegypti GN=BR-C PE=2 SV=1
CCG026397.1	1299	89	273	2.94403878	7.346889576	1.319336914	Up	0	0	AAEL002402-PA OS=Aedes aegypti GN=AAEL002402 PE=4 SV=1
CCG020100.1	4947	97	273	0.84254382	1.929171126	1.195157503	Up	1.90E-13	5.49E-13	AAEL012698-PA (Fragment) OS=Aedes aegypti GN=AAEL012698 PE=3 SV=1
CCG003907.1	3111	582	273	8.03869674	3.067698347	-1.389804998	Down	1.47E-42	6.64E-42	AAEL009536-PA OS=Aedes aegypti GN=AAEL009536 PE=4 SV=1
CCG003562.2	987	599	273	26.0778835	9.669310597	-1.431341848	Down	1.10E-45	5.04E-45	AAEL001308-PA OS=Aedes aegypti GN=AAEL001308 PE=4 SV=1
CCG020938.1	573	27	274	2.02475189	16.71652327	3.045457785	Up	2.28E-12	5.40E-12	AAEL000535-PA OS=Aedes aegypti GN=AAEL000535 PE=4 SV=1
CCG010979.1	2519	45	274	0.7676213	3.802527922	2.30849219	Up	0	0	AAEL008443-PA OS=Aedes aegypti GN=AAEL008443 PE=4 SV=1
CCG002611.1	933	86	274	3.96076868	10.26641783	1.374080532	Up	0	0	AAEL011988-PA OS=Aedes aegypti GN=AAEL011988 PE=4 SV=1
CCG022725.1	1896	93	274	2.10769268	5.051987255	1.261186476	Up	5.22E-14	1.75E-13	AAEL003859-PA (Fragment) OS=Aedes aegypti GN=AAEL003859 PE=4 SV=1
CCG027356.1	947	94	274	4.26521125	10.11464397	1.245756435	Up	0	0	AAEL012139-PA OS=Aedes aegypti GN=AAEL012139 PE=4 SV=1
CCG009196.1	3945	654	274	7.12349973	2.428027335	-1.552801539	Down	5.91E-56	2.82E-55	AAEL004000-PA OS=Aedes aegypti GN=TOLL10 PE=4 SV=1
CCG003703.1	1560	1355	274	37.3230708	6.140107587	-2.603731849	Down	2.87E-224	1.74E-223	AAEL003686-PA OS=Aedes aegypti GN=SRPN11 PE=3 SV=1

CCG028110.1	636	0	275	0.001	15.11560709	13.8837513	Up	4.44E-16	1.63E-15	Putative salivary secreted protein 62k-2 OS=Aedes albopictus PE=2 SV=1
CCG002658.1	768	2	275	0.11190035	12.51761213	6.805601012	Up	9.36E-05	0.00015402	AAEL009491-PA OS=Aedes aegypti GN=AAEL009491 PE=4 SV=1
CCG024116.1	533	4	275	0.32247456	18.03663436	5.805601012	Up	2.95E-05	5.04E-05	-
CCG022231.1	1371	4	275	0.12536757	7.012054057	5.805601012	Up	2.95E-05	5.03E-05	AAEL013582-PA OS=Aedes aegypti GN=AAEL013582 PE=4 SV=1
CCG007652.2	3033	33	275	0.46752431	3.169642635	2.761206893	Up	1.31E-13	4.01E-13	AAEL006034-PA OS=Aedes aegypti GN=AAEL006034 PE=4 SV=1
CCG008094.1	1620	49	275	1.29970185	5.934275378	2.190891168	Up	0	0	AAEL004232-PA OS=Aedes aegypti GN=AAEL004232 PE=4 SV=1
CCG005250.1	1437	50	275	1.49511951	6.689997295	2.161744822	Up	0	0	AAEL001925-PA OS=Aedes aegypti GN=AAEL001925 PE=4 SV=1
CCG018990.1	1488	58	275	1.67489557	6.460703033	1.947620017	Up	0	0	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018272 PE=4 SV=1
CCG027542.1	1998	64	275	1.37640792	4.811574631	1.805601012	Up	0	0	AAEL012916-PA OS=Aedes aegypti GN=AAEL012916 PE=4 SV=1
CCG017244.1	484	77	275	6.83609414	19.86265726	1.538814472	Up	0	0	AAEL000301-PA OS=Aedes aegypti GN=AAEL000301 PE=3 SV=1
CCG008808.1	461	87	275	8.10925577	20.85363582	1.362657516	Up	0	0	AAEL007241-PA OS=Aedes aegypti GN=AAEL007241 PE=4 SV=1
CCG000077.1	1305	101	275	3.32562697	7.366686676	1.147389529	Up	6.48E-13	1.65E-12	AGAP004026-PB OS=Anopheles gambiae GN=GPRMTH6 PE=4 SV=1
CCG004208.1	1083	108	275	4.28507049	8.876755413	1.05071351	Up	2.30E-11	5.03E-11	AAEL003244-PA OS=Aedes aegypti GN=AAEL003244 PE=4 SV=1
CCG006471.1	1734	703	275	17.4208324	5.544132706	-1.651779867	Down	1.58E-65	7.73E-65	AAEL011923-PA OS=Aedes aegypti GN=AAEL011923 PE=4 SV=1
CCG014399.1	624	856	275	58.9456616	15.40629185	-1.935865974	Down	9.81E-99	5.19E-98	AAEL008489-PA OS=Aedes aegypti GN=AAEL008489 PE=2 SV=1

CCG012761.1	1943	4	276	0.0884606	4.965766541	5.810837661	Up	2.95E-05	5.04E-05	ATP-binding cassette sub-family G member 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ019125 PE=3 SV=1
CCG019488.1	1249	42	276	1.44493903	7.724967485	2.418520238	Up	7.66E-14	2.47E-13	AAEL000600-PA OS=Aedes aegypti GN=AAEL006080 PE=3 SV=1
CCG014087.1	2562	78	276	1.30821206	3.765997029	1.525435442	Up	0	0	Single-minded OS=Culex quinquefasciatus GN=CpipJ_CPIJ009948 PE=4 SV=1
CCG001216.1	669	87	276	5.58799239	14.42224871	1.367894165	Up	0	0	AAEL004990-PA OS=Aedes aegypti GN=AAEL004990 PE=4 SV=1
CCG004402.1	3195	92	276	1.23731317	3.019869918	1.287275705	Up	1.75E-14	6.18E-14	DNA-repair protein complementing XP-G cells OS=Culex quinquefasciatus GN=CpipJ_CPIJ004184 PE=4 SV=1
CCG020908.1	1719	92	276	2.2997182	5.61284723	1.287275705	Up	1.75E-14	6.17E-14	AAEL014110-PA OS=Aedes aegypti GN=AAEL014110 PE=4 SV=1
CCG012729.1	1791	109	276	2.61513181	5.387205131	1.042653336	Up	2.82E-11	6.14E-11	AAEL000670-PA OS=Aedes aegypti GN=AAEL008870 PE=4 SV=1
CCG010044.1	474	8	277	0.72522759	20.42920394	4.81605537	Up	2.19E-06	3.96E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017096 PE=4 SV=1
CCG024517.1	1710	17	277	0.4271845	5.662831968	3.728592529	Up	3.39E-09	6.90E-09	AAEL004916-PA OS=Aedes aegypti GN=AAEL004916 PE=4 SV=1
CCG020302.1	1398	29	277	0.89136073	6.926639961	2.958074375	Up	6.66E-13	1.69E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000812 PE=4 SV=1
CCG010783.1	978	37	277	1.62564436	9.90127062	2.606602004	Up	5.17E-14	1.74E-13	Paralmyosin OS=Culex quinquefasciatus GN=CpipJ_CPIJ002902 PE=4 SV=1
CCG007291.1	1753	50	277	1.22560566	5.523926221	2.17219918	Up	0	0	AAEL007975-PA OS=Aedes aegypti GN=AAEL007975 PE=4 SV=1
CCG017558.1	705	70	277	4.26649847	13.73537967	1.686772353	Up	0	0	Hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1

CCG011006.1	627	89	277	6.09937222	15.44408719	1.340321939	Up	0	0	AAEL010170-PA OS=Aedes aegypti GN=AAEL010170 PE=3 SV=1
CCG006371.1	419	13	278	1.33318986	23.19427433	4.120814558	Up	6.34E-08	1.23E-07	-
CCG005235.1	1563	38	278	1.04468965	6.217786912	2.573326763	Up	4.26E-14	1.45E-13	AAEL003149-PA OS=Aedes aegypti GN=AAEL003149 PE=4 SV=1
CCG024789.1	1635	62	278	1.62943336	5.943976112	1.867057966	Up	0	0	AAEL011904-PA OS=Aedes aegypti GN=AAEL011904 PE=4 SV=1
CCG005391.2	1329	105	278	3.39490003	7.312566548	1.107008759	Up	2.17E-12	5.15E-12	AAEL006409-PA OS=Aedes aegypti GN=AAEL006409 PE=4 SV=1
CCG011609.1	1696	108	278	2.73628027	5.730189235	1.066366774	Up	1.02E-11	2.26E-11	AAEL013547-PA OS=Aedes aegypti GN=AAEL013547 PE=4 SV=1
CCG014670.1	3348	569	278	7.30280137	2.902748191	-1.331030566	Down	4.54E-39	2.02E-38	AAEL000121-PA (Fragment) OS=Aedes aegypti GN=AAEL000121 PE=4 SV=1
CCG013382.1	561	887	278	67.9396695	17.32335284	-1.971536018	Down	9.52E-105	5.07E-104	AAEL006451-PA OS=Aedes aegypti GN=AAEL006451 PE=4 SV=1
CCG023093.1	1248	900	278	30.9877894	7.787180243	-1.992526915	Down	9.50E-108	5.09E-107	AAEL007128-PA OS=Aedes aegypti GN=AAEL007128 PE=4 SV=1
CCG025474.1	1093	29	279	1.1400936	8.923475956	2.968453521	Up	6.66E-13	1.69E-12	AAEL004243-PA OS=Aedes aegypti GN=AAEL004243 PE=4 SV=1
CCG006579.2	1227	56	279	1.96112888	7.948948019	2.019079594	Up	0	0	AAEL012687-PA OS=Aedes aegypti GN=AAEL012687 PE=4 SV=1
CCG006630.1	1391	71	279	2.19327905	7.011760762	1.676687396	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000180 PE=4 SV=1
CCG027251.1	1173	74	279	2.71079315	8.314884245	1.61698115	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_16470 PE=4 SV=1
CCG020478.1	1056	75	279	3.05182774	9.236135624	1.597615825	Up	0	0	-
CCG006736.1	640	103	279	6.91544166	15.23962378	1.139933988	Up	5.82E-13	1.49E-12	AAEL000535-PA OS=Aedes aegypti GN=AAEL000535 PE=4 SV=1
CCG016181.1	1002	106	279	4.54570047	9.733891437	1.098514061	Up	2.85E-12	6.68E-12	GH regulated tbc protein-1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006102 PE=4 SV=1

CCG011510.1	728	574	279	33.8799831	13.39747146	-1.338472411	Down	1.00E-39	4.47E-39	AAEL007054-PA OS=Aedes aegypti GN=AAEL007054 PE=4 SV=1
CCG010280.1	1554	1435	279	39.6792594	6.276292934	-2.660400506	Down	3.12E-243	1.91E-242	AAEL002688-PA OS=Aedes aegypti GN=AAEL002688 PE=3 SV=1
CCG018537.1	2650	2099	279	34.0352728	3.680512913	-3.209051936	Down	0	0	AAEL008658-PA OS=Aedes aegypti GN=AAEL008658 PE=4 SV=1
CCG000159.1	1902	7	280	0.15814308	5.146328862	5.024241299	Up	4.31E-06	7.70E-06	AAEL002921-PA OS=Aedes aegypti GN=AAEL002921 PE=4 SV=1
CCG014037.1	753	41	280	2.33965354	12.99909362	2.474044216	Up	1.02E-13	3.19E-13	AAEL013188-PA OS=Aedes aegypti GN=AAEL013188 PE=4 SV=1
CCG001794.1	486	50	280	4.42075459	20.14057098	2.187740031	Up	0	0	AAEL002487-PA OS=Aedes aegypti GN=AAEL002487 PE=2 SV=1
CCG027865.1	906	83	280	3.93652094	10.80388245	1.456556789	Up	1.32E-13	4.01E-13	AAEL012387-PA OS=Aedes aegypti GN=AAEL012387 PE=4 SV=1
CCG010657.1	3735	87	280	1.00090145	2.620700802	1.388652725	Up	0	0	AAEL010619-PA OS=Aedes aegypti GN=AAEL010619 PE=3 SV=1
CCG020929.1	1752	101	280	2.47713653	5.58693921	1.173384738	Up	9.70E-14	3.05E-13	B250208117 KIK protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005754 PE=4 SV=1
CCG022766.1	2292	101	280	1.89351797	4.270644632	1.173384738	Up	9.70E-14	3.05E-13	AAEL010135-PA OS=Aedes aegypti GN=AAEL010135 PE=4 SV=1
CCG004074.1	437	1827	280	179.646923	22.39889587	-3.003664698	Down	0	0	-
CCG002133.1	11878	2952	280	10.6791258	0.824071182	-3.695880785	Down	0	0	AAEL001818-PA OS=Aedes aegypti GN=AAEL001818 PE=4 SV=1
CCG026777.1	420	16	281	1.63694227	23.38875184	3.836739524	Up	7.09E-09	1.42E-08	AAEL007250-PA OS=Aedes aegypti GN=AAEL007250 PE=4 SV=1
CCG025502.1	804	60	281	3.20669661	12.21800469	1.929848928	Up	0	0	AAEL012915-PA OS=Aedes aegypti GN=AAEL012915 PE=4 SV=1
CCG008403.1	1143	66	281	2.48119202	8.594292015	1.792345405	Up	0	0	UDP-galactose transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ007457 PE=4 SV=1
CCG007636.1	633	66	281	4.48025669	15.51860312	1.792345405	Up	0	0	AAEL000215-PA OS=Aedes aegypti GN=AAEL000215 PE=3 SV=1
CCG016847.1	1423	114	281	3.44241022	6.903215582	1.00384951	Up	7.50E-11	1.61E-10	AAEL008876-PA OS=Aedes aegypti GN=AAEL008876 PE=4 SV=1

CCG017331.1	18363	488	281	1.14192836	0.534949397	-1.093997814	Down	3.95E-25	1.61E-24	Amidophosphoribosyltransferase OS=Aedes aegypti GN=AAEL003581 PE=3 SV=1
CCG024358.1	2166	560	281	11.109442	4.535215038	-1.292543493	Down	7.83E-37	3.43E-36	AAEL014679-PA OS=Aedes aegypti GN=AAEL014679 PE=4 SV=1
CCG000528.1	2727	919	281	14.4808163	3.602228006	-2.007181527	Down	4.78E-111	2.57E-110	AAEL010294-PA OS=Aedes aegypti GN=AAEL010294 PE=4 SV=1
CCG025418.1	1182	10	282	0.36353413	8.340299535	4.519936461	Up	5.43E-07	1.01E-06	AAEL009465-PA OS=Aedes aegypti GN=AAEL009465 PE=4 SV=1
CCG016163.1	1356	49	282	1.55274115	7.270084108	2.227154712	Up	0	0	CpJ0B OS=Culex quinquefasciatus GN=CpipJ_CPIJ006746 PE=4 SV=1
CCG015495.1	588	59	282	4.31158902	16.76570417	1.959221507	Up	0	0	AAEL005789-PA OS=Aedes aegypti GN=AAEL005789 PE=4 SV=1
CCG022529.1	1095	76	282	2.98237427	9.00295347	1.593937043	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006253 PE=4 SV=1
CCG004122.1	1704	112	282	2.82430181	5.785348621	1.034509634	Up	2.31E-11	5.04E-11	AAEL007839-PA OS=Aedes aegypti GN=AAEL007839 PE=4 SV=1
CCG028246.1	318	543	282	73.3728487	31.00073601	-1.242943832	Down	9.93E-34	4.27E-33	AAEL013515-PA OS=Aedes aegypti GN=AAEL013515 PE=4 SV=1
CCG026404.1	660	21	283	1.36721883	14.98968534	3.454654024	Up	1.76E-10	3.73E-10	AAEL000672-PA OS=Aedes aegypti GN=AAEL000672 PE=3 SV=1
CCG016750.1	2351	26	283	0.47520761	4.208078403	3.146531729	Up	4.60E-12	1.05E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007173 PE=4 SV=1
CCG017566.1	1250	40	283	1.37503151	7.914553861	2.525043352	Up	6.86E-14	2.25E-13	AAEL006856-PA OS=Aedes aegypti GN=AAEL006856 PE=4 SV=1
CCG002852.1	1194	58	283	2.08730704	8.285755717	1.988990452	Up	0	0	AAEL002848-PA (Fragment) OS=Aedes aegypti GN=AAEL002848 PE=3 SV=1
CCG023515.1	2440	83	283	1.4616754	4.054587019	1.471932015	Up	1.32E-13	4.01E-13	AAEL007417-PA OS=Aedes aegypti GN=AAEL007417 PE=4 SV=1
CCG012893.1	1599	83	283	2.23044901	6.187112149	1.471932015	Up	1.32E-13	4.02E-13	AAEL009499-PA OS=Aedes aegypti GN=AAEL009499 PE=4 SV=1

CCG019690.1	861	95	283	4.74114377	11.49035113	1.277115838	Up	0	0	-
CCG012568.1	709	99	283	6.00000526	13.95372684	1.217614827	Up	1.42E-14	5.05E-14	AAEL002934-PA OS=Aedes aegypti GN=AAEL002934 PE=4 SV=1
CCG002897.1	2375	115	283	2.08063978	4.165554664	1.001481396	Up	6.99E-11	1.50E-10	AAEL009154-PA OS=Aedes aegypti GN=AAEL009154 PE=4 SV=1
CCG004226.1	1551	949	283	26.2916042	6.378589508	-2.04329283	Down	2.61E-117	1.42E-116	Abc transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ007889 PE=3 SV=1
CCG010332.1	897	38	284	1.8203455	11.06817236	2.60413281	Up	4.26E-14	1.45E-13	Methionine aminopeptidase OS=Aedes aegypti GN=AAEL013150 PE=3 SV=1
CCG007189.1	1638	41	284	1.07555502	6.061142004	2.494508319	Up	1.02E-13	3.19E-13	AAEL004874-PA OS=Aedes aegypti GN=AAEL004874 PE=4 SV=1
CCG023026.1	375	42	284	4.81261028	26.47506828	2.459742901	Up	7.66E-14	2.48E-13	AAEL005729-PA OS=Aedes aegypti GN=HOT_AEDAE PE=4 SV=1
CCG012279.1	540	65	284	5.17228287	18.38546408	1.82969251	Up	0	0	AAEL011200-PA OS=Aedes aegypti GN=AAEL011200 PE=4 SV=1
CCG012272.1	810	66	284	3.50123763	12.25697605	1.807666204	Up	0	0	AAEL012610-PA OS=Aedes aegypti GN=AAEL012610 PE=2 SV=1
CCG024800.1	1056	77	284	3.13320982	9.401657768	1.585273783	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_03220 PE=4 SV=1
CCG014690.1	978	86	284	3.77852472	10.15148323	1.425795569	Up	0	0	AAEL008636-PA OS=Aedes aegypti GN=AAEL008636 PE=4 SV=1
CCG020725.1	1027	96	284	4.0166451	9.667137881	1.267097823	Up	0	0	-
CCG014594.1	2099	107	284	2.19045336	4.729943117	1.110593337	Up	1.10E-12	2.69E-12	-
CCG007691.1	2732	522	284	8.21017623	3.634022915	-1.175845673	Down	7.03E-30	2.96E-29	AAEL012522-PA OS=Aedes aegypti GN=AAEL012522 PE=4 SV=1
CCG026430.1	801	524	284	28.1100386	12.39469489	-1.181362678	Down	3.35E-30	1.41E-29	AAEL005106-PA OS=Aedes aegypti GN=AAEL005106 PE=4 SV=1
CCG027317.1	2415	1262	284	22.4545777	4.111035447	-2.449435872	Down	1.01E-194	6.02E-194	AAEL010062-PA (Fragment) OS=Aedes aegypti GN=AAEL010062 PE=4 SV=1
CCG001950.1	579	0	285	0.001	17.20744193	14.07074502	Up	4.44E-16	1.66E-15	AAEL008835-PA OS=Aedes aegypti GN=AAEL008835 PE=4 SV=1

CCG007532.1	1350	0	285	0.001	7.380080652	12.84942087	Up	4.44E-16	1.63E-15	Tubulin alpha-2 chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ004534 PE=3 SV=1
CCG021444.1	1178	33	285	1.2037362	8.457647606	2.812737194	Up	1.31E-13	4.01E-13	Ribonuclease OS=Aedes aegypti GN=AAEL003510 PE=3 SV=1
CCG006768.1	1698	36	285	0.91101911	5.867555289	2.687206311	Up	6.13E-14	2.03E-13	AAEL003559-PA (Fragment) OS=Aedes aegypti GN=AAEL003559 PE=4 SV=1
CCG011163.1	1028	57	285	2.38256311	9.691740156	2.024241299	Up	0	0	-
CCG011015.1	828	59	285	3.06185307	12.03274019	1.974488264	Up	0	0	AAEL012523-PA OS=Aedes aegypti GN=AAEL012523 PE=4 SV=1
CCG019235.1	3054	73	285	1.02710892	3.26231463	1.667306754	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004892 PE=4 SV=1
CCG006059.1	597	90	285	6.47784944	16.68862459	1.365278217	Up	0	0	Tryptophan 2,3-dioxygenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ002864 PE=4 SV=1
CCG022221.1	663	93	285	6.02742884	15.02731354	1.317972502	Up	4.04E-14	1.38E-13	AAEL014296-PA OS=Aedes aegypti GN=AAEL014296 PE=4 SV=1
CCG021541.1	2324	996	285	18.4156005	4.287052014	-2.102870619	Down	1.11E-127	6.15E-127	AAEL002946-PA OS=Aedes aegypti GN=AAEL002946 PE=4 SV=1
CCG022586.1	2202	1464	285	28.5684339	4.524572607	-2.658568525	Down	6.45E-248	3.97E-247	AAEL008658-PA OS=Aedes aegypti GN=AAEL008658 PE=4 SV=1
CCG011855.1	971	21	286	0.92931455	10.2966706	3.469867118	Up	1.76E-10	3.72E-10	AAEL014754-PA OS=Aedes aegypti GN=AAEL014754 PE=3 SV=1
CCG009281.1	1367	37	286	1.16304329	7.313875023	2.652731175	Up	5.17E-14	1.74E-13	AAEL006298-PA (Fragment) OS=Aedes aegypti GN=AAEL006298 PE=4 SV=1
CCG021268.1	4524	49	286	0.46541048	2.210006003	2.247474696	Up	0	0	AAEL005047-PA OS=Aedes aegypti GN=AAEL005047 PE=4 SV=1
CCG025910.1	1711	52	286	1.30591829	5.843405702	2.161744822	Up	0	0	AAEL004163-PA OS=Aedes aegypti GN=AAEL004163 PE=4 SV=1
CCG000313.1	455	71	286	6.70516738	21.97377397	1.712437421	Up	0	0	AGAP007614-PA OS=Anopheles gambiae GN=AGAP007614 PE=4 SV=2

CCG008510.1	1239	80	286	2.77447843	8.069465018	1.540256446	Up	0	0	Aminomethyltransferase OS=Aedes aegypti GN=AAEL010276 PE=3 SV=1
CCG013998.1	711	85	286	5.13702875	14.06197912	1.452793604	Up	0	0	AAEL003756-PA OS=Aedes aegypti GN=AAEL003756 PE=4 SV=1
CCG007114.1	1008	111	286	4.73178625	9.918717418	1.067768674	Up	4.86E-12	1.11E-11	AAEL005208-PA OS=Aedes aegypti GN=AAEL005208 PE=2 SV=1
CCG028590.1	1812	574	286	13.6118254	5.517697106	-1.302722386	Down	3.72E-38	1.65E-37	AAEL013140-PA OS=Aedes aegypti GN=AAEL013140 PE=3 SV=1
CCG021828.1	2963	1010	286	14.6471252	3.374305487	-2.117955037	Down	1.22E-130	6.80E-130	AAEL009795-PA OS=Aedes aegypti GN=AAEL009795 PE=4 SV=1
CCG018321.1	920	4	287	0.18682493	10.90546243	5.86722013	Up	2.95E-05	5.04E-05	AAEL003648-PA OS=Aedes aegypti GN=AAEL003648 PE=4 SV=1
CCG025436.1	1062	23	287	0.93060631	9.447293252	3.343658174	Up	4.01E-11	8.68E-11	AAEL008938-PA OS=Aedes aegypti GN=AAEL008938 PE=4 SV=1
CCG023686.1	702	35	287	2.14236569	14.29205902	2.737937114	Up	6.39E-14	2.11E-13	AAEL014295-PA OS=Aedes aegypti GN=AAEL014295 PE=4 SV=1
CCG005520.1	325	38	287	5.02415359	30.87084749	2.619292617	Up	4.26E-14	1.45E-13	-
CCG021719.1	753	44	287	2.51084771	13.32407096	2.407788512	Up	3.91E-14	1.34E-13	Ribosome biogenesis protein (Fragment) OS=Aedes albopictus PE=2 SV=1
CCG026046.1	714	45	287	2.70817655	14.05185635	2.375367034	Up	0	0	AAEL003362-PA OS=Aedes aegypti GN=AAEL003362 PE=4 SV=1
CCG005857.1	1608	63	287	1.68351572	6.239443678	1.889940207	Up	0	0	AAEL011904-PA OS=Aedes aegypti GN=AAEL011904 PE=4 SV=1
CCG001488.1	2175	76	287	1.50147119	4.612885257	1.619292617	Up	0	0	AAEL011911-PA OS=Aedes aegypti GN=AAEL011911 PE=4 SV=1
CCG026536.1	681	92	287	5.80501554	14.73278331	1.343658174	Up	1.44E-14	5.11E-14	Pyridoxamine-phosphate oxidase OS=Aedes aegypti PE=2 SV=1
CCG017344.1	1362	101	287	3.18644875	7.366391655	1.209008648	Up	0	0	AAEL006769-PA OS=Aedes aegypti GN=AAEL006769 PE=3 SV=1
CCG018843.1	23442	574	287	1.05215543	0.427993577	-1.297686796	Down	6.18E-38	2.73E-37	AAEL006808-PA OS=Aedes aegypti GN=AAEL006808 PE=4 SV=1
CCG023923.1	869	824	287	40.7446045	11.54548381	-1.819280397	Down	1.76E-87	9.13E-87	AAEL012644-PA OS=Aedes aegypti GN=AAEL012644 PE=4 SV=1

CCG013769.1	852	1	288	0.05043396	11.81688229	7.872238205	Up	0.000587	0.00091865	Putative salivary mucin OS=Aedes albopictus PE=2 SV=1
CCG021994.1	651	71	288	4.68640731	15.46541277	1.722491086	Up	0	0	Hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG007647.1	1057	77	288	3.13024557	9.525055544	1.605451665	Up	0	0	III-proximal protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002301 PE=4 SV=1
CCG024926.1	732	112	288	6.5746042	13.75407611	1.064883283	Up	4.56E-12	1.05E-11	AAEL000129-PA OS=Aedes aegypti GN=AAEL006129 PE=4 SV=1
CCG017501.1	2603	494	288	8.15484015	3.867838536	-1.076129026	Down	8.72E-25	3.54E-24	AAEL001543-PA OS=Aedes aegypti GN=AAEL001543 PE=4 SV=1
CCG021263.1	6165	572	288	3.98681074	1.633087382	-1.287633132	Down	2.26E-37	9.95E-37	AAEL007759-PA OS=Aedes aegypti GN=AAEL007759 PE=4 SV=1
CCG025354.1	1231	598	288	20.8740059	8.178703258	-1.351763469	Down	6.33E-42	2.85E-41	AAEL008001-PA OS=Aedes aegypti GN=AAEL008001 PE=4 SV=1
CCG011044.1	16092	1299	288	3.46866053	0.625651486	-2.47094751	Down	1.89E-202	1.13E-201	AAEL009637-PA OS=Aedes aegypti GN=CAT-B3 PE=2 SV=1
CCG024717.1	2278	56	289	1.05632359	4.435005262	2.069883964	Up	0	0	AAEL009383-PA OS=Aedes aegypti GN=AAEL009383 PE=4 SV=1
CCG014835.1	1197	57	289	2.04617784	8.44021887	2.044348872	Up	0	0	AAEL011703-PA OS=Aedes aegypti GN=AAEL011703 PE=4 SV=1
CCG022436.1	1644	64	289	1.67278772	6.145341841	1.877238886	Up	0	0	AAEL007446-PA OS=Aedes aegypti GN=AAEL007446 PE=4 SV=1
CCG006734.1	1623	96	289	2.54164789	6.224856431	1.292276386	Up	0	0	AAEL006356-PA OS=Aedes aegypti GN=AAEL006356 PE=4 SV=1
CCG008677.1	730	105	289	6.18057827	13.83964656	1.162993369	Up	0	0	AAEL013010-PA OS=Aedes aegypti GN=AAEL013010 PE=4 SV=1
CCG002155.1	747	50	290	2.87615359	13.5714863	2.238366104	Up	0	0	AAEL001934-PA OS=Aedes aegypti GN=AAEL001934 PE=4 SV=1
CCG021417.1	998	60	290	2.58335078	10.1582167	1.975331698	Up	0	0	AAEL010970-PA OS=Aedes aegypti GN=AAEL010970 PE=4 SV=1
CCG018348.1	190	72	290	16.2832679	53.35736981	1.712297292	Up	0	0	-
CCG004960.1	519	497	290	41.1482815	19.53352652	-1.074879748	Down	6.98E-25	2.83E-24	AAEL010956-PA OS=Aedes aegypti GN=AAEL010956 PE=4 SV=1

CCG014686.1	906	504	290	23.9036934	11.18973539	-1.09505763	Down	6.14E-26	2.51E-25	AAEL006568-PA OS=Aedes aegypti GN=AAEL006568 PE=3 SV=1
CCG005370.1	827	21	291	1.0911299	12.30091722	3.494871124	Up	1.76E-10	3.72E-10	Alpha-tubulin (Fragment) OS=Cochliomyia hominivorax PE=2 SV=1
CCG020301.1	1236	38	291	1.32107598	8.230468075	2.639261033	Up	4.26E-14	1.45E-13	AAEL009454-PA OS=Aedes aegypti GN=AAEL009454 PE=4 SV=1
CCG003088.1	1462	44	291	1.29320679	6.958179576	2.427756928	Up	3.91E-14	1.34E-13	AAEL005132-PA OS=Aedes aegypti GN=AAEL005132 PE=4 SV=1
CCG003880.1	1065	51	291	2.0577056	9.55197985	2.214763205	Up	0	0	AAEL005019-PA OS=Aedes aegypti GN=AAEL005019 PE=4 SV=1
CCG020406.1	6393	69	291	0.4637747	1.591249576	1.77866409	Up	0	0	AAEL014742-PA (Fragment) OS=Aedes aegypti GN=AAEL014742 PE=4 SV=1
CCG015900.1	1193	82	291	2.95349391	8.527123672	1.529636542	Up	0	0	AAEL006694-PA OS=Aedes aegypti GN=AAEL006694 PE=4 SV=1
CCG004640.1	1786	95	291	2.28562418	5.695889441	1.317332938	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009123 PE=4 SV=1
CCG019786.1	826	1749	291	90.9855519	12.31580937	-2.885126027	Down	0	0	AAEL014889-PA (Fragment) OS=Aedes aegypti GN=AAEL014889 PE=2 SV=1
CCG007565.1	312	8115	291	1117.62627	32.60531584	-5.099186833	Down	0	0	AAEL001082-PA OS=Aedes aegypti GN=AAEL001082 PE=4 SV=1
CCG005940.1	579	0	292	0.001	17.63008086	14.10575147	Up	4.44E-16	1.66E-15	AAEL005005-PA OS=Aedes aegypti GN=AAEL005005 PE=4 SV=1
CCG022015.1	2431	13	292	0.22978468	4.19901967	4.191698045	Up	6.34E-08	1.23E-07	-
CCG000950.1	3357	35	292	0.4480014	3.04075568	2.762854746	Up	6.39E-14	2.10E-13	AAEL008178-PA OS=Aedes aegypti GN=AAEL008178 PE=3 SV=1
CCG016258.1	889	482	292	23.2974264	11.48235862	-1.020751574	Down	2.45E-22	9.74E-22	AAEL011967-PA OS=Aedes aegypti GN=AAEL011967 PE=4 SV=1
CCG012143.1	1842	745	292	17.3791815	5.541702941	-1.648958853	Down	3.62E-69	1.79E-68	AAEL005285-PA (Fragment) OS=Aedes aegypti GN=AAEL005285 PE=4 SV=1

CCG023583.1	2739	753	292	11.8131472	3.726840751	-1.664368292	Down	8.18E-71	4.06E-70	AAEL012655-PA OS=Aedes aegypti GN=AAEL012655 PE=3 SV=1
CCG025651.1	1317	820	292	26.7541248	7.750810036	-1.787342337	Down	5.59E-85	2.87E-84	AAEL002397-PA OS=Aedes aegypti GN=AAEL002397 PE=4 SV=1
CCG005588.1	1017	67	293	2.83084781	10.0715586	1.830980868	Up	0	0	AAEL012014-PA OS=Aedes aegypti GN=AAEL012014 PE=3 SV=1
CCG005769.1	1332	540	293	17.4201627	7.689771092	-1.179745539	Down	5.22E-31	2.21E-30	AAEL010988-PA (Fragment) OS=Aedes aegypti GN=AAEL010988 PE=4 SV=1
CCG020820.1	1699	556	293	14.0619026	6.028708119	-1.221871014	Down	1.28E-33	5.50E-33	-
CCG017107.1	1901	1536	293	34.7193647	5.388098419	-2.687892442	Down	3.23E-263	2.00E-262	AAEL007997-PA OS=Aedes aegypti GN=AAEL007997 PE=4 SV=1
CCG027863.1	1347	19	294	0.60610613	7.630091589	3.654058035	Up	7.73E-10	1.60E-09	AAEL012388-PA OS=Aedes aegypti GN=AAEL012388 PE=4 SV=1
CCG026300.1	1500	28	294	0.80210171	6.851822247	3.094630627	Up	1.17E-12	2.85E-12	AAEL013312-PA (Fragment) OS=Aedes aegypti GN=AAEL013312 PE=4 SV=1
CCG024512.1	2133	48	294	0.96697012	4.818440399	2.317023048	Up	5.35E-14	1.79E-13	AAEL004910-PA OS=Aedes aegypti GN=AAEL004910 PE=4 SV=1
CCG004212.1	1158	52	294	1.9295563	8.875417419	2.201545831	Up	0	0	AAEL011567-PA OS=Aedes aegypti GN=AAEL011567 PE=3 SV=1
CCG027333.1	631	67	294	4.56255502	16.28800851	1.835896358	Up	0	0	AAEL012135-PB OS=Aedes aegypti GN=GALE2 PE=4 SV=1
CCG015086.1	1085	73	294	2.89105127	9.472565319	1.71216099	Up	0	0	mediator complex OS=Culex quinquefasciatus GN=CpipJ_CPIJ013675 PE=4 SV=1
CCG014078.1	781	633	294	34.8269424	13.15970982	-1.404076141	Down	7.75E-47	3.57E-46	-
CCG000121.1	1049	660	294	27.0352954	9.79764859	-1.464336666	Down	7.35E-52	3.46E-51	AAEL006518-PA OS=Aedes aegypti GN=AAEL006518 PE=4 SV=1
CCG019298.1	1633	1137	294	29.9183027	6.293774263	-2.24903099	Down	2.24E-158	1.29E-157	AAEL001816-PA OS=Aedes aegypti GN=AAEL001816 PE=3 SV=1
CCG026452.1	450	0	295	0.001	22.91709255	14.4841364	Up	4.44E-16	1.64E-15	AAEL003185-PA OS=Aedes aegypti GN=AAEL003185 PE=4 SV=1
CCG027500.1	1100	1	295	0.0390634	9.375174225	7.906884348	Up	0.000587	0.00091683	AAEL002875-PA OS=Aedes aegypti GN=AAEL002875 PE=4 SV=1

CCG008913.1	1169	48	295	1.76436891	8.821806371	2.321921847	Up	5.35E-14	1.79E-13	AAEL011748-PA OS=Aedes aegypti GN=AAEL011748 PE=2 SV=1
CCG025973.1	663	74	295	4.79601864	15.5545877	1.697430982	Up	0	0	AAEL003817-PA OS=Aedes aegypti GN=AAEL003817 PE=2 SV=1
CCG020224.1	1596	92	295	2.47695212	6.461586246	1.383322392	Up	1.44E-14	5.12E-14	-
CCG014336.1	1248	759	295	26.1330357	8.263374718	-1.661071727	Down	3.65E-71	1.81E-70	AAEL011740-PA OS=Aedes aegypti GN=AAEL011740 PE=4 SV=1
CCG013856.1	1046	14	296	0.57512073	9.89259075	4.104411647	Up	3.07E-08	6.01E-08	AAEL006787-PA OS=Aedes aegypti GN=AAEL006787 PE=4 SV=1
CCG017159.1	3003	19	296	0.27186978	3.445770871	3.663839056	Up	7.73E-10	1.60E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001516 PE=4 SV=1
CCG019385.1	2352	21	296	0.38365834	4.399511022	3.519449147	Up	1.76E-10	3.72E-10	AAEL007753-PA OS=Aedes aegypti GN=AAEL007753 PE=4 SV=1
CCG012534.1	351	89	296	10.8954598	29.48048412	1.436033138	Up	0	0	AAEL012819-PA OS=Aedes aegypti GN=AAEL012819 PE=4 SV=1
CCG014699.1	861	974	296	48.6092004	12.01817645	-2.016011393	Down	2.94E-118	1.61E-117	AAEL010338-PA OS=Aedes aegypti GN=AAEL010338 PE=4 SV=1
CCG008602.1	1924	1102	296	24.6115632	5.378196427	-2.194141939	Down	6.81E-149	3.86E-148	AAEL012613-PA OS=Aedes aegypti GN=AAEL012613 PE=4 SV=1
CCG005828.1	1299	5	297	0.16539544	7.992769978	5.59470423	Up	1.59E-05	2.76E-05	AAEL006827-PA OS=Aedes aegypti GN=CYP12F8 PE=3 SV=1
CCG020736.1	1449	14	297	0.41516652	7.165361078	4.109277403	Up	3.07E-08	6.01E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014034 PE=4 SV=1
CCG023647.1	1992	39	297	0.84127492	5.212152712	2.631230106	Up	7.33E-14	2.38E-13	AAEL001468-PA OS=Aedes aegypti GN=AAEL001468 PE=4 SV=1
CCG027065.1	1392	40	297	1.23476249	7.45877026	2.59470423	Up	6.86E-14	2.24E-13	AAEL003178-PA OS=Aedes aegypti GN=AAEL003178 PE=4 SV=1
CCG012788.3	2118	92	297	1.86648517	4.902081304	1.393070369	Up	1.44E-14	5.11E-14	AAEL006023-PA OS=Aedes aegypti GN=AAEL006023 PE=4 SV=1
CCG009493.1	885	97	297	4.70967713	11.73176068	1.316719482	Up	3.62E-14	1.25E-13	AAEL001155-PA OS=Aedes aegypti GN=AAEL001155 PE=4 SV=1
CCG018157.1	489	102	297	8.96301213	21.23232761	1.244206983	Up	0	0	AAEL003029-PA OS=Aedes aegypti GN=AAEL003029 PE=4 SV=1

CCG000203.1	2208	116	297	2.25746794	4.702268207	1.058651329	Up	2.84E-12	6.67E-12	Zinc finger protein SZZA OS=Culex quinquefasciatus GN=CpipJ_CPIJ001554 PE=4 SV=1
CCG019367.2	2304	693	297	12.9244905	4.506340365	-1.520079218	Down	2.13E-57	1.02E-56	AAEL005181-PA OS=Aedes aegypti GN=AAEL005181 PE=4 SV=1
CCG023575.1	477	4751	297	427.985763	21.76647422	-4.297383167	Down	0	0	AAEL005768-PA OS=Aedes aegypti GN=AAEL005768 PE=4 SV=1
CCG001678.1	1279	8	298	0.26877082	8.14508716	4.921481724	Up	2.19E-06	3.97E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014034 PE=4 SV=1
CCG014170.1	1539	17	298	0.47464944	6.769049044	3.834018883	Up	3.39E-09	6.89E-09	AAEL002852-PA OS=Aedes aegypti GN=AAEL002852 PE=4 SV=1
CCG010770.1	246	30	298	5.24021154	42.34783121	3.014591129	Up	3.71E-13	9.94E-13	AAEL007539-PA OS=Aedes aegypti GN=AAEL007539 PE=4 SV=1
CCG011328.1	1143	37	298	1.39097129	9.114231389	2.712028359	Up	5.17E-14	1.74E-13	Flap endonuclease 1 OS=Anopheles darlingi GN=Fen1 PE=3 SV=1
CCG001300.1	1206	79	298	2.81476703	8.638114824	1.617700976	Up	0	0	AAEL006110-PA (Fragment) OS=Aedes aegypti GN=AAEL006110 PE=4 SV=1
CCG007708.1	3093	121	298	1.68100158	3.368110727	1.002618487	Up	2.10E-11	4.59E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_16340 PE=4 SV=1
CCG027596.1	1031	624	298	26.0069005	10.10433218	-1.363920495	Down	2.94E-44	1.34E-43	AAEL013871-PA OS=Aedes aegypti GN=AAEL013871 PE=4 SV=1
CCG022105.1	3342	718	298	9.23167847	3.117165314	-1.56635831	Down	5.03E-62	2.44E-61	AAEL012475-PA (Fragment) OS=Aedes aegypti GN=AAEL012475 PE=4 SV=1
CCG000417.1	1131	19	299	0.72186115	9.241843285	3.678387365	Up	7.73E-10	1.60E-09	AAEL003922-PA OS=Aedes aegypti GN=AAEL003922 PE=4 SV=1
CCG007091.1	1083	21	299	0.83320815	9.651454067	3.533997455	Up	1.76E-10	3.72E-10	Arginine N-methyltransferase 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003726 PE=4 SV=1
CCG026516.1	1659	45	299	1.16554434	6.30049714	2.434461782	Up	0	0	AAEL010013-PA OS=Aedes aegypti GN=AAEL010013 PE=4 SV=1

CCG022332.1	822	62	299	3.24102621	12.71596686	1.972118568	Up	0	0	AAEL011065-PA OS=Aedes aegypti GN=AAEL011065 PE=4 SV=1
CCG004641.1	1923	66	299	1.47478028	5.435530294	1.881920759	Up	0	0	AAEL007982-PA OS=Aedes aegypti GN=AAEL007982 PE=4 SV=1
CCG015382.1	678	679	299	43.0331118	15.41670318	-1.480952886	Down	3.47E-54	1.64E-53	AAEL001306-PA OS=Aedes aegypti GN=AAEL001306 PE=4 SV=1
CCG009651.2	1278	18	300	0.60520753	8.206168256	3.761206893	Up	1.62E-09	3.32E-09	Uncharacterized protein OS=Anopheles darlingi GN=AND_15689 PE=4 SV=1
CCG023156.1	936	22	300	1.00997239	11.20457589	3.471700276	Up	8.38E-11	1.80E-10	AAEL008124-PA OS=Aedes aegypti GN=AAEL008124 PE=4 SV=1
CCG012795.1	1398	42	300	1.29093623	7.501776131	2.538814472	Up	7.66E-14	2.47E-13	AAEL010332-PA OS=Aedes aegypti GN=AAEL010332 PE=4 SV=1
CCG021883.1	1219	45	300	1.58624943	8.603349493	2.439278798	Up	0	0	AAEL012240-PA (Fragment) OS=Aedes aegypti GN=AAEL012240 PE=4 SV=1
CCG026998.1	1695	52	300	1.31824555	6.187305623	2.230692176	Up	0	0	AAEL010498-PA OS=Aedes aegypti GN=AAEL010498 PE=4 SV=1
CCG010166.1	1075	54	300	2.15847969	9.755798169	2.176244392	Up	0	0	AAEL014712-PA OS=Aedes aegypti GN=AAEL014712 PE=4 SV=1
CCG025856.1	399	57	300	6.13853352	26.28441863	2.09824188	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001479 PE=4 SV=1
CCG022804.1	1083	62	300	2.45994787	9.683733178	1.976935584	Up	0	0	AAEL006311-PA OS=Aedes aegypti GN=AAEL006311 PE=4 SV=1
CCG017046.2	1011	73	300	3.10266135	10.3733759	1.741307335	Up	0	0	AAEL008794-PA OS=Aedes aegypti GN=AAEL008794 PE=4 SV=1
CCG025816.1	1905	78	300	1.75939071	5.505240437	1.645729675	Up	0	0	AAEL012156-PA OS=Aedes aegypti GN=AAEL012156 PE=4 SV=1
CCG003101.1	726	87	300	5.14926572	14.44556891	1.488188398	Up	0	0	AAEL002135-PA OS=Aedes aegypti GN=AAEL002135 PE=4 SV=1
CCG026069.1	1270	88	300	2.97743043	8.257860655	1.471700276	Up	2.93E-14	1.02E-13	AAEL005475-PA OS=Aedes aegypti GN=AAEL005475 PE=4 SV=1
CCG013716.1	1035	18	301	0.74729973	10.16660996	3.766007879	Up	1.62E-09	3.33E-09	AAEL006971-PA OS=Aedes aegypti GN=AAEL006971 PE=4 SV=1

CCG009352.1	774	21	301	1.16584551	13.59488541	3.543615458	Up	1.76E-10	3.73E-10	AAEL012675-PA OS=Aedes aegypti GN=AAEL012675 PE=4 SV=1
CCG011152.1	2745	39	301	0.61049896	3.833311952	2.650530662	Up	7.33E-14	2.38E-13	AAEL006631-PA OS=Aedes aegypti GN=AAEL006631 PE=4 SV=1
CCG023413.1	2541	46	301	0.77788579	4.141063089	2.412370925	Up	3.40E-14	1.18E-13	AAEL002994-PA OS=Aedes aegypti GN=AAEL002994 PE=4 SV=1
CCG004476.1	2381	48	301	0.86625252	4.419336963	2.35097038	Up	5.35E-14	1.79E-13	AAEL012727-PA OS=Aedes aegypti GN=AAEL012727 PE=4 SV=1
CCG022654.1	1301	55	301	1.81655296	8.087964111	2.154573167	Up	0	0	AAEL008602-PA OS=Aedes aegypti GN=AAEL008602 PE=4 SV=1
CCG026284.1	1640	66	301	1.72926981	6.416122749	1.891538761	Up	0	0	AAEL009246-PA OS=Aedes aegypti GN=AAEL009246 PE=3 SV=1
CCG003589.1	714	71	301	4.27290078	14.73731276	1.786185761	Up	0	0	AAEL006221-PA OS=Aedes aegypti GN=AAEL006221 PE=4 SV=1
CCG019393.1	702	85	301	5.20288809	14.98923263	1.526541944	Up	0	0	AAEL007351-PA OS=Aedes aegypti GN=AAEL007351 PE=4 SV=1
CCG021749.1	669	105	301	6.74412875	15.72861182	1.221687363	Up	0	0	AAEL002371-PA OS=Aedes aegypti GN=AAEL002371 PE=4 SV=1
CCG004909.1	456	664	301	62.5699644	23.07552919	-1.439106551	Down	7.44E-51	3.48E-50	AAEL005150-PA OS=Aedes aegypti GN=AAEL005150 PE=4 SV=1
CCG011433.1	2533	725	301	12.2988778	4.154141851	-1.565904304	Down	1.38E-62	6.71E-62	LINE-1 retrotransposon-like element, putative (Fragment) OS=Aspergillus flavus (strain ATCC 200026 / FGSC A1120 / NRRL 3357 / JCM 12722 / SRRC 167) GN=AFLA_063980 PE=4 SV=1
CCG018264.1	3825	814	301	9.14440888	2.750965048	-1.732952104	Down	7.62E-81	3.88E-80	AAEL009195-PA OS=Aedes aegypti GN=AAEL009195 PE=4 SV=1
CCG025048.1	558	100	302	7.70066929	18.92007094	1.296861753	Up	0	0	AAEL013472-PA OS=Aedes aegypti GN=AAEL013472 PE=4 SV=1
CCG024035.1	823	595	302	31.065604	12.82794603	-1.276027915	Down	2.78E-38	1.23E-37	AAEL005666-PA OS=Aedes aegypti GN=AAEL005666 PE=4 SV=1
CCG022568.1	1596	916	302	24.6618276	6.614912021	-1.898485845	Down	8.29E-103	4.41E-102	AAEL000311-PA OS=Aedes aegypti GN=AAEL000311 PE=4 SV=1

CCG027334.1	1538	1117	302	31.2075381	6.864369041	-2.184695527	Down	4.46E-150	2.53E-149	AGAP000380-PA OS=Anopheles gambiae GN=AgaP_AGAP000380 PE=3 SV=5
CCG022384.1	1686	1709	302	43.5559172	6.261802838	-2.798218739	Down	6.00E-306	3.80E-305	AAEL011831-PA OS=Aedes aegypti GN=AAEL011831 PE=4 SV=1
CCG015626.1	1522	32	303	0.90343726	6.959499252	2.945487187	Up	1.77E-13	5.17E-13	AAEL007864-PA OS=Aedes aegypti GN=AAEL007864 PE=4 SV=1
CCG004429.1	1281	94	303	3.15312651	8.268819564	1.390898336	Up	0	0	AAEL001600-PA OS=Aedes aegypti GN=AAEL001600 PE=4 SV=1
CCG000944.1	4764	2836	303	25.5798	2.223416848	-3.52415463	Down	0	0	AAEL008160-PA OS=Aedes aegypti GN=AAEL008160 PE=4 SV=1
CCG014859.1	456	7	304	0.65962312	23.30551785	5.142885795	Up	4.31E-06	7.70E-06	AAEL008463-PA OS=Aedes aegypti GN=AAEL008463 PE=3 SV=1
CCG015554.1	1752	14	304	0.34336546	6.065819714	4.142885795	Up	3.07E-08	6.00E-08	AAEL005452-PA OS=Aedes aegypti GN=AAEL005452 PE=4 SV=1
CCG023580.1	939	33	304	1.51011847	11.31769557	2.905846598	Up	1.31E-13	4.00E-13	AAEL006853-PA OS=Aedes aegypti GN=AAEL006853 PE=4 SV=1
CCG025562.2	974	48	304	2.11760499	10.9110022	2.365278217	Up	5.35E-14	1.79E-13	AAEL012747-PA OS=Aedes aegypti GN=AAEL012747 PE=4 SV=1
CCG009130.1	639	48	304	3.22777349	16.63116767	2.365278217	Up	5.35E-14	1.80E-13	-
CCG017116.1	1356	64	304	2.02807007	7.83725379	1.950240717	Up	0	0	AAEL000477-PA OS=Aedes aegypti GN=AAEL000477 PE=4 SV=1
CCG017838.1	699	65	304	3.99575501	15.20359963	1.927872904	Up	0	0	NAD(P)+H ₂ O-hydrate epimerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ004022 PE=3 SV=1
CCG028451.1	1636	88	304	2.31133047	6.49591451	1.490809099	Up	2.93E-14	1.02E-13	AAEL008648-PA OS=Aedes aegypti GN=AAEL008648 PE=4 SV=1
CCG011240.1	1713	520	304	13.0439358	6.203920688	-1.072127096	Down	7.40E-26	3.03E-25	AAEL008596-PA OS=Aedes aegypti GN=SPZ3A PE=4 SV=1
CCG019696.1	685	2081	304	130.540172	15.51433013	-3.072820532	Down	0	0	AAEL006829-PA OS=Aedes aegypti GN=AAEL006829 PE=4 SV=1
CCG014945.1	477	2133	304	192.147681	22.27948876	-3.108427533	Down	0	0	AAEL003662-PA OS=Aedes aegypti GN=AAEL003662 PE=4 SV=1
CCG015046.1	597	2312	304	166.408754	17.80119956	-3.224684965	Down	0	0	AAEL011070-PA OS=Aedes aegypti GN=CTLGA3 PE=2 SV=1

CCG007987.1	1056	105	305	4.27255884	10.09685077	1.240733119	Up	0	0	AAEL004974-PA OS=Aedes aegypti GN=AAEL004974 PE=4 SV=1
CCG024908.1	2271	771	305	14.5881398	4.694968919	-1.635608414	Down	1.06E-70	5.26E-70	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011195 PE=4 SV=1
CCG015867.1	1667	0	306	0.001	6.417056204	12.6476959	Up	4.44E-16	1.64E-15	AAEL000123-PA OS=Aedes aegypti GN=AAEL006123 PE=3 SV=1
CCG010399.1	957	16	306	0.71840727	11.1778816	3.959701047	Up	7.09E-09	1.42E-08	AAEL011648-PA OS=Aedes aegypti GN=AAEL011648 PE=3 SV=1
CCG002035.1	1097	74	306	2.8985965	9.751351588	1.750247681	Up	0	0	AAEL001322-PA OS=Aedes aegypti GN=AAEL001322 PE=4 SV=1
CCG025414.1	2185	79	306	1.55359681	4.895758669	1.655920298	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009399 PE=4 SV=1
CCG022646.1	2787	104	306	1.60346337	3.838260744	1.259261328	Up	0	0	AAEL006397-PA (Fragment) OS=Aedes aegypti GN=AAEL006397 PE=4 SV=1
CCG015343.1	1983	724	306	15.6883953	5.394469336	-1.540144841	Down	4.97E-61	2.41E-60	AAEL003316-PA OS=Aedes aegypti GN=AAEL003316 PE=4 SV=1
CCG023257.1	399	74	307	7.96932421	26.89772173	1.754954684	Up	0	0	AAEL012078-PA OS=Aedes aegypti GN=AAEL012078 PE=4 SV=1
CCG022582.1	855	76	307	3.81953197	12.55227014	1.716480536	Up	0	0	AAEL006915-PA OS=Aedes aegypti GN=AAEL006915 PE=4 SV=1
CCG021448.1	1908	76	307	1.71158272	5.624838034	1.716480536	Up	0	0	AAEL003546-PA OS=Aedes aegypti GN=AAEL003546 PE=4 SV=1
CCG014515.1	1202	95	307	3.39611047	8.928611455	1.394552441	Up	0	0	AAEL014651-PA OS=Aedes aegypti GN=AAEL014651 PE=4 SV=1
CCG008606.1	3660	98	307	1.15055574	2.932292615	1.349698205	Up	0	0	Putative uncharacterized protein OS=Drosophila simulans GN=pol PE=4 SV=1
CCG010178.1	1376	103	307	3.21648449	7.79955739	1.277907522	Up	0	0	AAEL006463-PA OS=Aedes aegypti GN=AAEL006463 PE=4 SV=1
CCG000515.1	2397	7	308	0.12548525	4.491927095	5.161744822	Up	4.31E-06	7.70E-06	AAEL008530-PA OS=Aedes aegypti GN=AAEL008530 PE=4 SV=1

CCG007690.1	1752	27	308	0.66220481	6.145633131	3.214212242	Up	2.28E-12	5.40E-12	AAEL001426-PA OS=Aedes aegypti GN=AAEL001426 PE=4 SV=1
CCG010823.1	504	33	308	2.81349453	21.36339136	2.924705625	Up	1.31E-13	4.01E-13	AAEL0008801-PA OS=Aedes aegypti GN=AAEL008801 PE=4 SV=1
CCG010311.1	501	40	308	3.43071733	21.49131586	2.64717165	Up	6.86E-14	2.25E-13	AAEL0007250-PA OS=Aedes aegypti GN=AAEL007250 PE=4 SV=1
CCG022069.1	1311	75	308	2.4582228	8.212928487	1.740281054	Up	0	0	AAEL010524-PA OS=Aedes aegypti GN=AAEL010524 PE=4 SV=1
CCG015767.2	1944	114	308	2.51983012	5.538657019	1.13620973	Up	1.33E-13	4.02E-13	AGAP008245-PA (Fragment) OS=Anopheles gambiae GN=AGAP008245 PE=4 SV=3
CCG025483.1	1746	723	308	17.7933093	6.166752145	-1.528752092	Down	2.68E-60	1.29E-59	AAEL000838-PA OS=Aedes aegypti GN=AAEL000838 PE=4 SV=1
CCG003542.1	643	773	308	51.6572393	16.74517768	-1.625224859	Down	3.04E-70	1.51E-69	AAEL002074-PA OS=Aedes aegypti GN=AAEL002074 PE=4 SV=1
CCG020003.2	1118	9	309	0.34591021	9.661992417	4.80385123	Up	1.09E-06	2.01E-06	-
CCG008373.1	795	27	309	1.45934948	13.58755663	3.21888873	Up	2.28E-12	5.41E-12	AAEL005009-PA OS=Aedes aegypti GN=AAEL005009 PE=4 SV=1
CCG006541.1	693	77	309	4.77441496	15.58745674	1.706989691	Up	0	0	-
CCG028053.1	2640	78	309	1.26956034	4.091707395	1.688374013	Up	0	0	AAEL002820-PA OS=Aedes aegypti GN=AAEL002820 PE=4 SV=1
CCG015227.1	609	121	309	8.53750064	17.73745078	1.054912994	Up	1.10E-12	2.71E-12	AAEL012658-PA OS=Aedes aegypti GN=AAEL012658 PE=4 SV=1
CCG002508.1	1638	806	309	21.1438377	6.594693237	-1.680859797	Down	1.00E-76	5.04E-76	AAEL000101-PA OS=Aedes aegypti GN=AAEL000101 PE=4 SV=1
CCG019377.1	954	2	310	0.0900833	11.35960776	6.978437609	Up	9.36E-05	0.00015382	AAEL000643-PA OS=Aedes aegypti GN=AAEL006643 PE=4 SV=1
CCG012580.1	1587	24	310	0.64982585	6.828648897	3.393475108	Up	1.93E-11	4.23E-11	AAEL003700-PA OS=Aedes aegypti GN=AAEL003700 PE=4 SV=1
CCG016263.1	954	30	310	1.35124952	11.35960776	3.071547013	Up	3.71E-13	9.95E-13	-
CCG008060.1	917	49	310	2.29609269	11.81795616	2.363727765	Up	0	0	AAEL008171-PA OS=Aedes aegypti GN=AAEL008171 PE=4 SV=1
CCG024387.1	1904	70	310	1.57976965	5.691736239	1.849154592	Up	0	0	AAEL002739-PA OS=Aedes aegypti GN=AAEL002739 PE=4 SV=1

CCG001923.1	1087	98	310	3.87399631	9.969701747	1.363727765	Up	0	0	AAEL001573-PA OS=Aedes aegypti GN=AAEL001573 PE=4 SV=1
CCG002172.1	1881	109	310	2.49000589	5.761332163	1.210253284	Up	0	0	AAEL011366-PA OS=Aedes aegypti GN=AAEL011366 PE=4 SV=1
CCG012609.1	1527	4	311	0.11255988	7.119858596	5.983083974	Up	2.95E-05	5.03E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005075 PE=4 SV=1
CCG018427.1	996	9	311	0.38828073	10.91568682	4.813158973	Up	1.09E-06	2.01E-06	AAEL006547-PA OS=Aedes aegypti GN=AAEL006547 PE=2 SV=1
CCG011683.1	1725	16	311	0.39855986	6.302622653	3.983083974	Up	7.09E-09	1.42E-08	AAEL000053-PA OS=Aedes aegypti GN=AAEL000053 PE=4 SV=1
CCG022454.1	525	41	311	3.35573166	20.70861729	2.625531969	Up	1.02E-13	3.19E-13	Structural maintenance of chromosomes protein OS=Aedes aegypti GN=AAEL001655 PE=3 SV=1
CCG020881.1	2319	51	311	0.94500063	4.688238066	2.310658632	Up	0	0	AAEL010663-PA OS=Aedes aegypti GN=AAEL010663 PE=4 SV=1
CCG017421.1	588	56	311	4.09235568	18.48983686	2.175729052	Up	0	0	AAEL012081-PA OS=Aedes aegypti GN=AAEL012081 PE=4 SV=1
CCG024154.1	582	60	311	4.42986955	18.68045374	2.076193378	Up	0	0	AAEL011841-PA OS=Aedes aegypti GN=AAEL011841 PE=3 SV=1
CCG020245.1	618	62	311	4.31087952	17.59227197	2.028887664	Up	0	0	AAEL003288-PA OS=Aedes aegypti GN=AAEL003288 PE=3 SV=1
CCG019452.1	4631	62	311	0.5752804	2.347662292	2.028887664	Up	0	0	AAEL003737-PA (Fragment) OS=Aedes aegypti GN=AAEL003737 PE=4 SV=1
CCG016027.1	3141	65	311	0.88921769	3.461325717	1.960716161	Up	0	0	AAEL011173-PA OS=Aedes aegypti GN=AAEL011173 PE=4 SV=1
CCG025863.1	896	86	311	4.12432721	12.13395544	1.556819219	Up	0	0	-
CCG004687.1	888	90	311	4.35504067	12.24327036	1.491230878	Up	0	0	AAEL004867-PA OS=Aedes aegypti GN=AAEL004867 PE=4 SV=1
CCG016524.1	501	725	311	62.1817517	21.70064686	-1.518753211	Down	6.87E-60	3.31E-59	AAEL003467-PA OS=Aedes aegypti GN=AAEL003467 PE=4 SV=1
CCG021610.1	1275	54	312	1.81989464	8.554495963	2.232827921	Up	0	0	AAEL008393-PA OS=Aedes aegypti GN=AAEL008393 PE=4 SV=1

CCG018752.2	3892	75	312	0.82803959	2.802410677	1.758896732	Up	0	0	AAEL006864-PA OS=Aedes aegypti GN=AAEL006864 PE=4 SV=1
CCG024948.1	584	81	312	5.95984333	18.67633965	1.64786542	Up	4.46E-14	1.51E-13	-
CCG020167.1	2360	118	312	2.14848673	4.621602692	1.105072373	Up	4.60E-14	1.55E-13	AAEL006622-PA OS=Aedes aegypti GN=AAEL006622 PE=3 SV=1
CCG015143.1	2728	606	312	9.54532961	3.998160687	-1.255458561	Down	5.11E-38	2.26E-37	AAEL014428-PA OS=Aedes aegypti GN=AAEL014428 PE=3 SV=1
CCG013932.1	1773	1193	312	28.9130814	6.151710295	-2.232662905	Down	1.41E-164	8.13E-164	Acetylcholinesterase OS=Aedes albopictus GN=AP-AChE PE=2 SV=1
CCG010906.1	417	2	313	0.20608985	26.23966578	6.992332051	Up	9.36E-05	0.00015364	Histone H2B OS=Aedes aegypti GN=AAEL015674 PE=3 SV=1
CCG023769.1	653	55	313	3.61919664	16.7564175	2.210972337	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009802 PE=4 SV=1
CCG012002.1	927	91	313	4.21817244	11.8036037	1.484537411	Up	0	0	AAEL004900-PA OS=Aedes aegypti GN=AAEL004900 PE=3 SV=1
CCG007261.1	1243	120	313	4.14832514	8.802848455	1.085441455	Up	1.18E-13	3.65E-13	AAEL011762-PA OS=Aedes aegypti GN=AAEL011762 PE=4 SV=1
CCG024472.1	2244	1431	313	27.4018227	4.876087625	-2.490475906	Down	6.88E-225	4.17E-224	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005148 PE=4 SV=1
CCG015870.1	848	2527	313	128.047782	12.90323187	-3.310877898	Down	0	0	NADH dehydrogenase OS=Ochlerotatus taeniorhynchus PE=2 SV=1
CCG004463.1	645	11	314	0.73281718	17.01844792	4.537502334	Up	2.67E-07	5.03E-07	AAEL009356-PA OS=Aedes aegypti GN=AAEL009356 PE=4 SV=1
CCG017508.1	1125	39	314	1.48961747	9.757243472	2.711531734	Up	7.33E-14	2.38E-13	AAEL000786-PA OS=Aedes aegypti GN=AAEL000786 PE=4 SV=1
CCG011631.1	1680	43	314	1.09982059	6.533868397	2.570669198	Up	8.46E-14	2.70E-13	AAEL009924-PA OS=Aedes aegypti GN=AAEL009924 PE=4 SV=1
CCG000111.2	1185	54	314	1.95811449	9.263205828	2.242046451	Up	0	0	AAEL012236-PA OS=Aedes aegypti GN=AAEL012236 PE=4 SV=1
CCG003731.1	4922	68	314	0.59364932	2.23017044	1.909471111	Up	0	0	AAEL012505-PA OS=Aedes aegypti GN=AAEL012505 PE=4 SV=1

CCG009573.1	909	104	314	4.91622926	12.07579638	1.296494235	Up	0	0	AAEL009293-PA OS=Aedes aegypti GN=AAEL009293 PE=4 SV=1
CCG011918.1	1908	529	314	11.9135166	5.75309167	-1.050189959	Down	1.88E-25	7.66E-25	AAEL010268-PA OS=Aedes aegypti GN=AAEL010268 PE=4 SV=1
CCG004185.1	1059	651	314	26.4148227	10.36534363	-1.34957978	Down	2.23E-45	1.02E-44	AAEL012518-PA OS=Aedes aegypti GN=AAEL012518 PE=4 SV=1
CCG001401.1	1157	1596	314	59.2737221	9.487380213	-2.643310984	Down	2.39E-268	1.49E-267	AAEL002971-PA OS=Aedes aegypti GN=AAEL002971 PE=4 SV=1
CCG012514.1	1050	0	315	0.001	10.48748303	13.35638086	Up	4.44E-16	1.65E-15	AAEL007783-PA OS=Aedes aegypti GN=AAEL007783 PE=4 SV=1
CCG017875.1	1644	26	315	0.67957001	6.698209965	3.301081504	Up	4.60E-12	1.05E-11	AAEL010362-PA OS=Aedes aegypti GN=AAEL010362 PE=4 SV=1
CCG014942.1	971	28	315	1.23908606	11.3407386	3.1941663	Up	1.17E-12	2.86E-12	AAEL001358-PA OS=Aedes aegypti GN=AAEL001358 PE=4 SV=1
CCG008178.1	624	40	315	2.75447017	17.64720702	2.679593127	Up	6.86E-14	2.25E-13	Transmembrane trafficking protein OS=Aedes aegypti PE=2 SV=1
CCG021103.1	2042	56	315	1.17840604	5.392682264	2.1941663	Up	0	0	AAEL001986-PA OS=Aedes aegypti GN=AAEL001986 PE=3 SV=1
CCG002443.1	1521	84	315	2.37308199	7.239879805	1.609203799	Up	0	0	AAEL011053-PA OS=Aedes aegypti GN=AAEL011053 PE=4 SV=1
CCG000820.1	1617	110	315	2.9231112	6.810053917	1.220161509	Up	0	0	AAEL006805-PA OS=Aedes aegypti GN=CYP9J2 PE=3 SV=2
CCG017061.1	2703	971	315	15.4360386	4.073939025	-1.921806263	Down	1.15E-110	6.20E-110	AAEL009530-PA OS=Aedes aegypti GN=AAEL009530 PE=4 SV=1
CCG015079.1	498	4	316	0.34513843	22.18236036	6.006093952	Up	2.95E-05	5.04E-05	AAEL012441-PA OS=Aedes aegypti GN=AAEL012441 PE=4 SV=1
CCG007878.1	390	7	316	0.77125165	28.32516785	5.19873903	Up	4.31E-06	7.69E-06	AAEL004909-PA OS=Aedes aegypti GN=AAEL004909 PE=4 SV=1
CCG022620.1	1002	42	316	1.8011266	11.02476593	2.613776529	Up	7.66E-14	2.47E-13	-
CCG022673.1	411	47	316	4.91381393	26.8778965	2.4515051	Up	0	0	Histone H3 OS=Gecko japonicus PE=2 SV=1
CCG017795.1	498	55	316	4.74565342	22.18236036	2.224734238	Up	0	0	Nuclear receptor FTZ-F1 protein OS=Aedes aegypti GN=FTZ-F1 PE=2 SV=1

CCG020264.1	399	76	316	8.18471135	27.68625429	1.758166439	Up	0	0	AAEL008355-PA OS=Aedes aegypti GN=AAEL008355 PE=4 SV=1
CCG008921.1	1422	102	316	3.08221725	7.768505949	1.33366861	Up	0	0	AAEL012917-PA (Fragment) OS=Aedes aegypti GN=AAEL012917 PE=4 SV=1
CCG027457.1	1605	119	316	3.18591802	6.882751065	1.111276189	Up	0	0	AAEL001144-PA OS=Aedes aegypti GN=AAEL001144 PE=4 SV=1
CCG021954.1	1029	825	316	34.4509534	10.73548636	-1.682156357	Down	1.41E-78	7.14E-78	AAEL008227-PA OS=Aedes aegypti GN=AAEL008227 PE=4 SV=1
CCG014168.1	1760	30	317	0.73243866	6.29646235	3.103761638	Up	3.71E-13	9.95E-13	AAEL002849-PA OS=Aedes aegypti GN=AAEL002849 PE=4 SV=1
CCG011611.2	1893	49	317	1.11226466	5.854080157	2.39594239	Up	0	0	AAEL013796-PA OS=Aedes aegypti GN=AAEL013796 PE=4 SV=1
CCG012492.1	1068	116	317	4.66712473	10.37619264	1.152671239	Up	1.26E-13	3.88E-13	AAEL004448-PA OS=Aedes aegypti GN=AAEL004448 PE=4 SV=1
CCG027218.1	852	1879	317	94.7654124	13.00677669	-2.865097117	Down	0	0	AAEL008601-PA OS=Aedes aegypti GN=AAEL008601 PE=2 SV=1
CCG005313.1	1047	2645	317	108.552959	10.58431111	-3.358399773	Down	0	0	AAEL003067-PB OS=Aedes aegypti GN=AAEL003067 PE=4 SV=1
CCG023636.1	1611	26	318	0.69349044	6.900516458	3.314756441	Up	4.60E-12	1.05E-11	AAEL007931-PA OS=Aedes aegypti GN=AAEL007931 PE=4 SV=1
CCG005525.1	12363	56	318	0.19463764	0.899193724	2.207841237	Up	0	0	Cytoplasmic dynein heavy chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ005887 PE=4 SV=1
CCG025806.1	2025	102	318	2.16440145	5.489744204	1.342770817	Up	0	0	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG008667.1	1600	120	318	3.2227301	6.947957508	1.108305563	Up	0	0	AAEL011053-PA OS=Aedes aegypti GN=AAEL011053 PE=4 SV=1
CCG025552.1	507	2234	318	189.338042	21.92649312	-3.110217311	Down	0	0	AAEL008006-PA OS=Aedes aegypti GN=AAEL008006 PE=4 SV=1
CCG020385.1	768	50	319	2.79750876	14.52043007	2.375869628	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013304 PE=4 SV=1
CCG023007.1	750	87	319	4.98448921	14.86892039	1.576782322	Up	0	0	AAEL011338-PA OS=Aedes aegypti GN=AAEL011338 PE=4 SV=1

CCG026473.1	394	0	320	0.001	28.39250905	14.79322273	Up	4.44E-16	1.66E-15	Chromosome undetermined SCAF14565, whole genome shotgun sequence OS=Tetraodon nigroviridis GN=GSTENG00016696001 PE=3 SV=1
CCG017120.1	759	15	320	0.84920424	14.73866741	4.117350703	Up	1.48E-08	2.93E-08	AAEL009090-FA OS=Aedes aegypti GN=AAEL009090 PE=4 SV=1
CCG024425.1	1164	32	320	1.18129855	9.610522824	3.024241299	Up	1.77E-13	5.16E-13	Ribonuclease OS=Aedes aegypti GN=AAEL003510 PE=3 SV=1
CCG017979.1	957	35	320	1.57151589	11.68928795	2.894958282	Up	6.39E-14	2.11E-13	Aldo/keto reductase OS=Aedes aegypti PE=2 SV=1
CCG008813.1	902	77	320	3.66814808	12.40204941	1.757454758	Up	0	0	AAEL007238-FA OS=Aedes aegypti GN=AAEL007238 PE=4 SV=1
CCG022599.1	1306	89	320	2.9282591	8.565580832	1.548507868	Up	0	0	Serine/threonine protein kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ011728 PE=4 SV=1
CCG009971.1	12781	111	320	0.37318211	0.875256128	1.229825432	Up	0	0	-
CCG021715.1	2295	1290	320	24.1529227	4.874356674	-2.308914052	Down	2.45E-185	1.44E-184	AAEL007028-FA OS=Aedes aegypti GN=AAEL007028 PE=4 SV=1
CCG011440.1	577	26	321	1.93624454	19.4481921	3.328302973	Up	4.60E-12	1.05E-11	-
CCG000374.1	2772	26	321	0.40303503	4.048198717	3.328302973	Up	4.60E-12	1.05E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014679 PE=4 SV=1
CCG025943.1	435	38	321	3.75367797	25.79679734	2.780815177	Up	4.26E-14	1.45E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_12728 PE=4 SV=1
CCG003800.1	1023	83	321	3.486303	10.96931265	1.65370326	Up	1.32E-13	4.02E-13	AAEL001041-FA OS=Aedes aegypti GN=AAEL001041 PE=4 SV=1
CCG024002.1	1257	123	321	4.2046757	8.927292636	1.086228186	Up	0	0	AAEL008822-FA OS=Aedes aegypti GN=AAEL008822 PE=4 SV=1
CCG000773.1	498	557	321	48.0605265	22.53334708	-1.092790826	Down	2.10E-28	8.78E-28	S95 ribosomal protein L35, mitochondrial OS=Culex quinquefasciatus GN=CpipJ_CPIJ019136 PE=4 SV=1

CCG004903.1	11847	1174	321	4.25816396	0.947210842	-2.168474002	Down	3.70E-156	2.12E-155	AAEL007293-PA OS=Aedes aegypti GN=AAEL007293 PE=2 SV=1
CCG022471.1	1527	2615	321	73.5860223	7.348792956	-3.32385254	Down	0	0	Dihydrolipoyl dehydrogenase OS=Aedes aegypti GN=AAEL006928 PE=3 SV=1
CCG023511.1	718	26	322	1.5560071	15.6776673	3.332790364	Up	4.60E-12	1.05E-11	AAEL000215-PA OS=Aedes aegypti GN=AAEL000215 PE=3 SV=1
CCG014195.1	1689	76	322	1.93351085	6.66463299	1.785302568	Up	0	0	AAEL012057-PA OS=Aedes aegypti GN=AAEL012057 PE=4 SV=1
CCG013816.1	669	641	322	41.1713003	16.82595683	-1.290950465	Down	7.77E-42	3.49E-41	AAEL013528-PA OS=Aedes aegypti GN=TPX1 PE=4 SV=1
CCG023555.1	2539	1273	322	21.5441009	4.43346401	-2.280786622	Down	3.47E-180	2.03E-179	AAEL001499-PA OS=Aedes aegypti GN=AAEL001499 PE=4 SV=1
CCG009484.1	603	27	323	1.92401797	18.72557777	3.282816056	Up	2.28E-12	5.41E-12	AAEL004976-PA OS=Aedes aegypti GN=AAEL004976 PE=4 SV=1
CCG004524.1	1749	34	323	0.83531788	6.45598822	2.950240717	Up	8.82E-14	2.80E-13	AGAP005314-PA OS=Anopheles gambiae GN=AGAP005314 PE=4 SV=2
CCG008296.1	3942	82	323	0.89384024	2.864414865	1.680151554	Up	0	0	AAEL006614-PA OS=Aedes aegypti GN=AAEL006614 PE=4 SV=1
CCG013239.1	1368	86	323	2.70131373	8.254037571	1.611438804	Up	0	0	AGAP006454-PA OS=Anopheles gambiae GN=AGAP006454 PE=4 SV=4
CCG023037.1	219	89	323	17.4625862	51.55946757	1.561970128	Up	0	0	AAEL002578-PA OS=Aedes aegypti GN=AAEL002578 PE=4 SV=1
CCG015268.1	1863	117	323	2.69858237	6.060935801	1.167338839	Up	9.73E-14	3.06E-13	AAEL006244-PA OS=Aedes aegypti GN=AAEL006244 PE=4 SV=1
CCG012524.1	2063	603	323	12.5597431	5.473351138	-1.198310633	Down	2.93E-35	1.27E-34	AAEL002579-PA OS=Aedes aegypti GN=AAEL002579 PE=4 SV=1
CCG018736.1	738	697	323	40.5825271	15.30016721	-1.407311287	Down	1.43E-51	6.73E-51	AAEL010075-PA OS=Aedes aegypti GN=AAEL010075 PE=3 SV=1
CCG017086.1	3509	790	323	9.67400694	3.217875006	-1.588005285	Down	2.04E-69	1.01E-68	AAEL015018-PA (Fragment) OS=Aedes aegypti GN=AAEL015018 PE=4 SV=1

CCG008353.1	861	8	324	0.39925421	13.15503098	5.042163207	Up	2.19E-06	3.97E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018760 PE=4 SV=1
CCG004511.1	1470	15	324	0.43846668	7.705089574	4.135272611	Up	1.48E-08	2.93E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014034 PE=4 SV=1
CCG000370.1	2193	23	324	0.45066297	5.164834325	3.518601251	Up	4.01E-11	8.67E-11	AAEL003691-PA OS=Aedes aegypti GN=AAEL003691 PE=4 SV=1
CCG021558.1	987	39	324	1.69789225	11.47566532	2.756760988	Up	7.33E-14	2.38E-13	AAEL008320-PA OS=Aedes aegypti GN=AAEL008320 PE=4 SV=1
CCG009574.1	1455	98	324	2.89418144	7.784523487	1.427453363	Up	0	0	AAEL009282-PA OS=Aedes aegypti GN=AAEL009282 PE=4 SV=1
CCG006593.1	1179	699	324	25.4756951	9.606854686	-1.406985439	Down	1.07E-51	5.04E-51	Cytosolic acid phosphatase OS=Culex quinquefasciatus GN=CpipJ_CPIJ008635 PE=4 SV=1
CCG017341.1	3039	1007	324	14.2384083	3.727042341	-1.933684761	Down	1.20E-115	6.53E-115	AAEL003025-PA OS=Aedes aegypti GN=AAEL003025 PE=4 SV=1
CCG023926.1	813	5587	324	295.291399	13.93171178	-4.405694898	Down	0	0	AAEL012646-PA OS=Aedes aegypti GN=AAEL012646 PE=4 SV=1
CCG002306.1	576	25	325	1.86500584	19.72472214	3.402752922	Up	9.32E-12	2.08E-11	AAEL010504-PA OS=Aedes aegypti GN=AAEL010504 PE=4 SV=1
CCG026907.1	1260	29	325	0.98898596	9.017015834	3.188628117	Up	6.66E-13	1.69E-12	AAEL003744-PA (Fragment) OS=Aedes aegypti GN=AAEL003744 PE=4 SV=1
CCG012407.1	460	36	325	3.3628488	24.6987825	2.87668411	Up	6.13E-14	2.03E-13	AAEL009759-PA OS=Aedes aegypti GN=AAEL009759 PE=4 SV=1
CCG006333.1	2628	60	325	0.98104417	4.32322677	2.139718516	Up	0	0	Uncharacterized protein OS=Nomascus leucogenys GN=ZNF808 PE=4 SV=1
CCG008877.1	921	61	325	2.84598677	12.33598257	2.115871774	Up	1.75E-14	6.18E-14	AAEL009222-PA OS=Aedes aegypti GN=AAEL009222 PE=4 SV=1
CCG008401.1	1170	88	325	3.23191166	9.710632437	1.587177493	Up	2.93E-14	1.02E-13	AAEL008120-PA OS=Aedes aegypti GN=AAEL008120 PE=3 SV=1

CCG010037.1	253	119	325	20.2110609	44.90687728	1.151791348	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_05744 PE=4 SV=1
CCG013104.1	540	893	325	71.0592093	21.03970361	-1.755907253	Down	3.83E-90	1.99E-89	AAEL004729-FA OS=Aedes aegypti GN=AAEL004729 PE=4 SV=1
CCG023784.1	4026	982	325	10.4809437	2.822016878	-1.892970103	Down	1.19E-109	6.41E-109	AAEL012395-FA OS=Aedes aegypti GN=AAEL012395 PE=3 SV=1
CCG026029.1	1367	2144	325	67.3936438	8.311221617	-3.019480079	Down	0	0	Larval chymotrypsin-like protein (Precursor) OS=Aedes aegypti PE=2 SV=1
CCG018556.1	906	40	326	1.89711853	12.57880599	2.729113263	Up	6.86E-14	2.25E-13	AAEL002179-FA OS=Aedes aegypti GN=AAEL002179 PE=4 SV=1
CCG012315.1	2652	62	326	1.00457147	4.2972844	2.096845048	Up	0	0	AAEL013690-FA OS=Aedes aegypti GN=AAEL013690 PE=4 SV=1
CCG012731.1	1233	70	326	2.43948209	9.242820947	1.921758341	Up	0	0	AAEL008867-FA OS=Aedes aegypti GN=AAEL008867 PE=4 SV=1
CCG007749.1	459	92	326	8.61267012	24.82875431	1.527479402	Up	1.44E-14	5.12E-14	Nuclear protein localization 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006953 PE=4 SV=1
CCG020117.1	1689	611	326	15.5444096	6.747423462	-1.203987212	Down	5.80E-36	2.53E-35	AAEL012981-FA OS=Aedes aegypti GN=AAEL012981 PE=4 SV=1
CCG016857.1	1488	653	326	18.8570139	7.658869777	-1.299897823	Down	4.96E-43	2.24E-42	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011712 PE=4 SV=1
CCG017570.1	984	25	327	1.09171074	11.61723222	3.41160384	Up	9.32E-12	2.08E-11	Trypsin-like salivary secreted protein OS=Aedes albopictus PE=2 SV=1
CCG010565.1	1550	49	327	1.35839806	7.375068713	2.440750185	Up	0	0	AAEL011655-FA OS=Aedes aegypti GN=AAEL011655 PE=3 SV=1
CCG015011.1	1365	54	327	1.69990159	8.37462015	2.300572527	Up	0	0	AAEL010118-FA OS=Aedes aegypti GN=AAEL010118 PE=4 SV=1
CCG010526.1	1770	111	327	2.69471217	6.458393505	1.261044163	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017830 PE=4 SV=1
CCG025860.1	3537	674	327	8.18818239	3.231935681	-1.341144752	Down	1.75E-46	8.03E-46	AAEL000712-FA OS=Aedes aegypti GN=AAEL000712 PE=4 SV=1

CCG027708.4	4228	1007	327	10.2342769	2.703726704	-1.920387939	Down	1.39E-114	7.51E-114	AAEL004582-PA OS=Aedes aegypti GN=AAEL004582 PE=4 SV=1
CCG026865.1	1575	1422	327	38.7955318	7.25800413	-2.41824572	Down	7.59E-216	4.57E-215	Putative adenosine deaminase OS=Aedes albopictus PE=2 SV=1
CCG020712.1	2156	1693	327	33.742004	5.302113407	-2.669906229	Down	9.45E-288	5.91E-287	AAEL007191-PA OS=Aedes aegypti GN=AAEL007191 PE=2 SV=1
CCG006433.1	222	5923	327	1146.44026	51.49259687	-4.476652341	Down	0	0	AAEL009337-PA OS=Aedes aegypti GN=AAEL009337 PE=3 SV=1
CCG014329.1	1008	45	328	1.91829172	11.37531228	2.568012112	Up	0	0	AAEL013887-PA OS=Aedes aegypti GN=AAEL013887 PE=4 SV=1
CCG009733.1	1710	45	328	1.13078249	6.705447241	2.568012112	Up	0	0	AAEL010176-PA OS=Aedes aegypti GN=AAEL010176 PE=4 SV=1
CCG026859.1	2078	76	328	1.57155911	5.517957065	1.811937695	Up	0	0	Eukaryotic translation initiation factor 4 gamma 3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009979 PE=4 SV=1
CCG002803.1	1834	132	328	3.09269628	6.252080033	1.015471089	Up	1.36E-12	3.30E-12	AAEL000602-PA (Fragment) OS=Aedes aegypti GN=GPRNNB2 PE=4 SV=2
CCG019192.1	1425	33	329	0.99508859	8.071068813	3.019862858	Up	1.31E-13	4.01E-13	tetracycline resistance OS=Culex quinquefasciatus GN=CpipJ_CPIJ003321 PE=4 SV=1
CCG005236.1	663	60	329	3.88866377	17.34731985	2.157366382	Up	0	0	AAEL010173-PA OS=Aedes aegypti GN=AAEL010173 PE=4 SV=1
CCG018902.1	4623	63	329	0.58557069	2.487837564	2.086977054	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010464 PE=4 SV=1
CCG022230.1	1335	78	329	2.51059124	8.615185811	1.778854759	Up	0	0	AAEL013099-PA OS=Aedes aegypti GN=AAEL013099 PE=4 SV=1
CCG027928.1	291	26	330	3.83922027	39.64340665	3.3681957	Up	4.60E-12	1.05E-11	AAEL005232-PA OS=Aedes aegypti GN=AAEL005232 PE=4 SV=1
CCG025857.1	1230	34	330	1.18778128	9.379049866	2.981172577	Up	8.82E-14	2.80E-13	AAEL000699-PA OS=Aedes aegypti GN=AAEL000699 PE=4 SV=1
CCG014836.1	1347	34	330	1.08461097	8.564388519	2.981172577	Up	8.82E-14	2.79E-13	AAEL011705-PA OS=Aedes aegypti GN=AAEL011705 PE=4 SV=1

CCG024516.1	1569	73	330	1.99922921	7.352601233	1.878810859	Up	0	0	AAEL004895-PA OS=Aedes aegypti GN=AAEL004895 PE=4 SV-1
CCG005255.1	1932	76	330	1.69032082	5.971134231	1.820707905	Up	0	0	AAEL012747-PA OS=Aedes aegypti GN=AAEL012747 PE=4 SV-1
CCG001440.1	984	96	330	4.19216923	11.72381233	1.483672917	Up	0	0	AAEL003240-PA OS=Aedes aegypti GN=AAEL003246 PE=4 SV-1
CCG020713.1	420	1023	330	104.661996	27.46721746	-1.929955012	Down	3.76E-117	2.05E-116	AAEL001429-PA OS=Aedes aegypti GN=AAEL001429 PE=4 SV-1
CCG022969.1	2214	6	331	0.11644915	5.226372905	5.48803811	Up	8.38E-06	1.47E-05	AAEL014020-PA OS=Aedes aegypti GN=AAEL014020 PE=4 SV-1
CCG019685.1	1464	34	331	0.997931	7.903818041	2.985537769	Up	8.82E-14	2.79E-13	AAEL005592-PA OS=Aedes aegypti GN=AAEL005592 PE=4 SV-1
CCG009434.1	768	60	331	3.35701052	15.06665314	2.166110015	Up	0	0	AAEL014183-PA OS=Aedes aegypti GN=AAEL014183 PE=4 SV-1
CCG014059.1	13534	8307	331	26.3742859	0.854971894	-4.947111228	Down	0	0	AAEL008032-PA OS=Aedes aegypti GN=AAEL008032 PE=4 SV-1
CCG012610.1	1347	20	332	0.63800645	8.616293904	3.75542454	Up	3.69E-10	7.72E-10	AAEL008419-PA OS=Aedes aegypti GN=AAEL008419 PE=4 SV-1
CCG021159.1	2129	33	332	0.66604098	5.451455091	3.032958516	Up	1.31E-13	4.01E-13	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV-1
CCG015262.1	948	35	332	1.58643535	12.24277203	2.948069618	Up	6.39E-14	2.10E-13	AAEL004960-PA OS=Aedes aegypti GN=AAEL004960 PE=4 SV-1
CCG005846.1	1048	43	332	1.76307117	11.0745686	2.65108788	Up	8.46E-14	2.70E-13	AAEL003786-PA OS=Aedes aegypti GN=AAEL003786 PE=4 SV-1
CCG027633.1	997	72	332	3.10313028	11.6410711	1.907427634	Up	0	0	AAEL003390-PA OS=Aedes aegypti GN=AAEL003390 PE=4 SV-1
CCG013794.1	1202	103	332	3.68209872	9.655697079	1.390852108	Up	0	0	G173-specific cyprin-C OS=Culex quinquefasciatus GN=CpipJ_CPIJ006239 PE=3 SV-1
CCG005829.1	2652	3041	332	49.2726105	4.376375524	-3.492977466	Down	0	0	AAEL006822-PA OS=Aedes aegypti GN=AAEL006822 PE=4 SV-1
CCG007048.1	606	8	333	0.56725722	19.20974615	5.081691571	Up	2.19E-06	3.96E-06	AAEL001624-PA OS=Aedes aegypti GN=AAEL001624 PE=4 SV-1

CCG006231.1	2883	55	333	0.81974867	4.037844664	2.300331857	Up	0	0	isoleducyl-tRNA synthetase OS=Culex quinquefasciatus GN=CpipJ_CPIJ015134 PE=4 SV=1
CCG018080.1	1968	57	333	1.24455024	5.915196222	2.248801557	Up	0	0	isoleducyl-tRNA synthetase OS=Culex quinquefasciatus GN=CpipJ_CPIJ015134 PE=4 SV=1
CCG005707.1	1665	65	333	1.67749715	6.991655354	2.059323758	Up	0	0	isoleducyl-tRNA synthetase OS=Aedes aegypti GN=AAEL002744 PE=4 SV=1
CCG023663.1	732	70	333	4.10912763	15.9031505	1.952408554	Up	0	0	Eukaryotic translation initiation factor 6 OS=Aedes aegypti GN=EIF6 PE=3 SV=1
CCG020543.1	786	75	333	4.10016552	14.81056764	1.85287288	Up	0	0	AAEL013168-PA (Fragment) OS=Aedes aegypti GN=AAEL013168 PE=3 SV=1
CCG003273.1	537	79	333	6.32143209	21.67803755	1.777910823	Up	0	0	AAEL007902-PA OS=Aedes aegypti GN=AAEL007902 PE=4 SV=1
CCG012958.1	1482	82	333	2.37754267	7.854997412	1.724139566	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003369 PE=4 SV=1
CCG004941.1	1929	82	333	1.82660354	6.034788059	1.724139566	Up	0	0	AAEL013943-PB (Fragment) OS=Aedes aegypti GN=AAEL013943 PE=4 SV=1
CCG013289.1	1161	95	333	3.51604202	10.02679256	1.511835963	Up	0	0	AAEL011356-PA OS=Aedes aegypti GN=AAEL011356 PE=4 SV=1
CCG028064.1	1338	98	333	3.14726008	8.7003783	1.466981727	Up	0	0	AAEL000905-PA (Fragment) OS=Aedes aegypti GN=AAEL000905 PE=4 SV=2
CCG014697.1	987	663	333	28.8641682	11.7944338	-1.291173489	Down	3.18E-43	1.44E-42	AAEL010337-PB OS=Aedes aegypti GN=AAEL010337 PE=4 SV=1
CCG028152.1	1308	1053	333	34.5926075	8.899928261	-1.95859815	Down	6.41E-123	3.53E-122	AAEL012888-PA (Fragment) OS=Aedes aegypti GN=AAEL012888 PE=4 SV=1
CCG018029.3	3213	1284	333	17.1718454	3.623126724	-2.244737916	Down	3.55E-178	2.08E-177	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000250 PE=4 SV=1

CCG015355.1	2037	1387	333	29.2582336	5.714828751	-2.356060501	Down	4.32E-204	2.59E-203	AAEL011280-PA (Fragment) OS=Aedes aegypti GN=AAEL011280 PE=3 SV=1
CCG011394.1	516	1569	333	130.657972	22.56028327	-2.533938066	Down	2.11E-251	1.30E-250	AAEL000663-PA OS=Aedes aegypti GN=AAEL000663 PE=4 SV=1
CCG012576.1	975	1632	333	71.924725	11.93959607	-2.590733771	Down	3.59E-268	2.23E-267	AAEL014452-PA OS=Aedes aegypti GN=AAEL014452 PE=3 SV=1
CCG027729.1	1565	30	334	0.82370098	7.460744052	3.179126901	Up	3.71E-13	9.94E-13	AAEL013254-PA OS=Aedes aegypti GN=AAEL013254 PE=4 SV=1
CCG023102.1	1104	31	334	1.20657769	10.57614533	3.131821186	Up	2.40E-13	6.78E-13	AAEL001472-PA OS=Aedes aegypti GN=AAEL001472 PE=3 SV=1
CCG009886.1	2520	54	334	0.92078003	4.633358906	2.331129994	Up	0	0	AAEL000491-PA OS=Aedes aegypti GN=AAEL006491 PE=4 SV=1
CCG005917.1	2325	82	334	1.51549172	5.021963201	1.728465492	Up	0	0	AAEL014992-PA (Fragment) OS=Aedes aegypti GN=AAEL014992 PE=4 SV=1
CCG007524.1	993	84	334	3.63490202	11.75837305	1.693700074	Up	0	0	AAEL000290-PA OS=Aedes aegypti GN=AAEL000290 PE=4 SV=1
CCG014902.1	961	1462	334	65.37123	12.14991097	-2.4277101	Down	6.75E-223	4.09E-222	AAEL012764-PA OS=Aedes aegypti GN=AAEL012764 PE=3 SV=1
CCG017410.1	957	6	335	0.26940272	12.23722332	5.505367988	Up	8.38E-06	1.47E-05	-
CCG024207.1	643	9	335	0.60144263	18.2130991	4.920405488	Up	1.09E-06	2.01E-06	AAEL010085-PA OS=Aedes aegypti GN=AAEL010085 PE=4 SV=1
CCG002550.1	1077	37	335	1.47621187	10.8737444	2.880877124	Up	5.17E-14	1.74E-13	AAEL006788-PA (Fragment) OS=Aedes aegypti GN=AAEL006788 PE=2 SV=1
CCG019865.1	367	56	335	6.55668975	31.91014365	2.282975567	Up	0	0	AAEL003435-PA OS=Aedes aegypti GN=AAEL003435 PE=4 SV=1
CCG008858.1	1044	101	335	4.15703371	11.21745471	1.432119006	Up	0	0	AAEL008416-PA OS=Aedes aegypti GN=AAEL008416 PE=4 SV=1
CCG015317.1	3168	128	335	1.73615089	3.696661212	1.090330489	Up	6.93E-14	2.26E-13	AAEL009926-PA OS=Aedes aegypti GN=AAEL009926 PE=4 SV=1
CCG026662.1	1371	135	335	4.23115549	8.541956761	1.013514892	Up	9.52E-13	2.36E-12	Undn OS=Culex quinquefasciatus GN=Cuq11_CUL002280 PE=4

CCG013364.1	3114	639	335	8.81748889	3.760765163	-1.229341632	Down	9.65E-39	4.28E-38	Angiotensin-converting enzyme OS=Culex quinquefasciatus GN=CpipJ_CPIJ004086 PE=4 SV=1
CCG025981.1	2037	86	336	1.81413705	5.766313694	1.668365872	Up	0	0	AAEL001966-PA OS=Aedes aegypti GN=AAEL001966 PE=4 SV=1
CCG021038.1	579	98	336	7.27294299	20.28666839	1.479920782	Up	0	0	AAEL002008-PA (Fragment) OS=Aedes aegypti GN=AAEL002008 PE=4 SV=1
CCG021520.1	492	99	336	8.64634904	23.87394511	1.465274007	Up	0	0	AAEL005355-PA OS=Aedes aegypti GN=AAEL005355 PE=4 SV=1
CCG016368.1	3810	5054	336	56.9997477	3.082934644	-4.208579322	Down	0	0	AAEL013228-PA (Fragment) OS=Aedes aegypti GN=AAEL013228 PE=4 SV=1
CCG004458.1	1743	26	337	0.64097137	6.759001304	3.398478267	Up	4.60E-12	1.05E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000812 PE=4 SV=1
CCG010505.1	2151	44	337	0.87897179	5.476959215	2.639486366	Up	3.91E-14	1.34E-13	AAEL009531-PA OS=Aedes aegypti GN=AAEL009531 PE=4 SV=1
CCG004806.1	770	90	337	5.02243651	15.29992113	1.607064889	Up	0	0	AAEL014183-PA OS=Aedes aegypti GN=AAEL014183 PE=4 SV=1
CCG019961.1	870	132	337	6.51954594	13.54130951	1.054523866	Up	2.58E-13	7.23E-13	AAEL011912-PA OS=Aedes aegypti GN=AAEL011912 PE=2 SV=1
CCG021265.1	1818	832	337	19.664917	6.480164616	-1.601521733	Down	6.68E-74	3.35E-73	AAEL006216-PA OS=Aedes aegypti GN=AAEL006216 PE=4 SV=1
CCG012583.1	1611	44	338	1.17359921	7.334511204	2.643761021	Up	3.91E-14	1.35E-13	AAEL003657-PA OS=Aedes aegypti GN=AAEL003657 PE=4 SV=1
CCG022674.1	1035	55	338	2.28341585	11.41632613	2.321832927	Up	0	0	Cytosine-specific methyltransferase OS=Aedes aegypti GN=AAEL006166 PE=3 SV=1
CCG020842.1	1360	65	338	2.05370055	8.688159963	2.080824827	Up	0	0	AAEL002069-PA OS=Aedes aegypti GN=AAEL002069 PE=4 SV=1
CCG019978.1	1449	82	338	2.43168961	8.154518667	1.745640635	Up	0	0	AAEL013478-PA OS=Aedes aegypti GN=AAEL013478 PE=4 SV=1
CCG007644.1	234	114	338	20.9339733	50.49528867	1.270302626	Up	8.48E-14	2.70E-13	AAEL001247-PA OS=Aedes aegypti GN=AAEL001247 PE=4 SV=1

CCG022291.1	600	137	338	9.81142274	19.69316258	1.005160557	Up	1.15E-12	2.81E-12	AAEL009140-PA OS=Aedes aegypti GN=AAEL009140 PE=4 SV=1
CCG005692.2	195	620	338	136.62172	60.59434641	-1.172931765	Down	4.83E-35	2.09E-34	AAEL002081-PA OS=Aedes aegypti GN=AAEL002081 PE=4 SV=1
CCG024918.1	1962	656	338	14.3670468	6.022373878	-1.254359365	Down	5.85E-41	2.62E-40	AAEL006778-PA (Fragment) OS=Aedes aegypti GN=GPRFZ2 PE=4 SV=2
CCG016030.1	894	42	339	2.01871236	13.25599086	2.715137244	Up	7.66E-14	2.47E-13	AAEL015062-PA OS=Aedes aegypti GN=AAEL015062 PE=4 SV=1
CCG009496.1	972	52	339	2.29879239	12.1922385	2.407014949	Up	0	0	AAEL001185-PA OS=Aedes aegypti GN=AAEL001185 PE=4 SV=1
CCG005145.1	1895	84	339	1.90472702	6.253749776	1.715137244	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012958 PE=4 SV=1
CCG018552.1	965	120	339	5.34338669	12.28067961	1.200564071	Up	0	0	Putative uncharacterized protein OS=Aedes aegypti PE=2 SV=1
CCG019768.1	1872	584	339	13.4050881	6.330585377	-1.082369892	Down	2.90E-29	1.22E-28	AAEL005491-PA OS=Aedes aegypti GN=AAEL005491 PE=3 SV=1
CCG014293.1	1339	823	339	26.4108227	8.850527129	-1.577293953	Down	1.52E-71	7.58E-71	AAEL012021-PA OS=Aedes aegypti GN=AAEL012021 PE=4 SV=1
CCG025187.1	1245	20	340	0.69027686	9.546838637	3.789776045	Up	3.69E-10	7.71E-10	-
CCG013487.1	1455	74	340	2.18540231	8.1689444	1.902250774	Up	0	0	AAEL014654-PA OS=Aedes aegypti GN=AAEL014654 PE=4 SV=1
CCG001368.1	823	85	340	4.43794343	14.44205845	1.702313204	Up	0	0	AAEL004276-PA OS=Aedes aegypti GN=AAEL004276 PE=4 SV=1
CCG004749.1	1383	108	340	3.35555411	8.594225671	1.356816638	Up	0	0	AAEL009798-PA OS=Aedes aegypti GN=AAEL009798 PE=4 SV=1
CCG022100.1	1488	658	340	19.0014015	7.987778295	-1.250239634	Down	7.11E-41	3.19E-40	AAEL007773-PA OS=Aedes aegypti GN=AAEL007773 PE=4 SV=1
CCG021933.1	625	1076	340	73.9766951	19.01730256	-1.959758223	Down	1.16E-125	6.41E-125	AAEL000467-PA OS=Aedes aegypti GN=AAEL000467 PE=2 SV=1
CCG008561.1	1954	43	341	0.94559805	6.100702344	2.689676378	Up	8.46E-14	2.70E-13	AAEL003672-PA OS=Aedes aegypti GN=AAEL003672 PE=4 SV=1
CCG016228.1	501	63	341	5.4033798	23.79395685	2.138661209	Up	0	0	AAEL004879-PA OS=Aedes aegypti GN=AAEL004879 PE=4 SV=1

CCG007263.1	1071	73	341	2.92884279	11.13050642	1.926116574	Up	0	0	AAEL012418-PA OS=Aedes aegypti GN=AAEL012418 PE=4 SV=1
CCG021643.1	666	81	341	5.2260488	17.89905763	1.77609113	Up	4.46E-14	1.51E-13	AAEL003678-PA OS=Aedes aegypti GN=AAEL003678 PE=3 SV=1
CCG025055.1	699	91	341	5.59405701	17.05403774	1.608146493	Up	0	0	AAEL005586-PA OS=Aedes aegypti GN=AAEL005586 PE=4 SV=1
CCG026495.1	1059	718	341	29.1333989	11.25663114	-1.371898901	Down	4.20E-51	1.97E-50	AAEL011597-PA OS=Aedes aegypti GN=AAEL011597 PE=4 SV=1
CCG027472.1	670	1255	341	80.488085	17.79219758	-2.177530516	Down	1.11E-167	6.44E-167	AAEL008490-PA OS=Aedes aegypti GN=AAEL008490 PE=4 SV=1
CCG015538.1	2777	8	342	0.1237875	4.305268511	5.120165719	Up	2.19E-06	3.96E-06	AGAP001039-PB OS=Anopheles gambiae GN=AgaP_AGAP001039 PE=3 SV=1
CCG002364.1	1250	12	342	0.41250945	9.564584525	4.535203218	Up	1.30E-07	2.49E-07	AAEL006080-PA OS=Aedes aegypti GN=AAEL006080 PE=3 SV=1
CCG022032.1	615	17	342	1.18778128	19.44021245	4.032702877	Up	3.39E-09	6.88E-09	AAEL004861-PA OS=Aedes aegypti GN=AAEL004861 PE=4 SV=1
CCG007522.1	1257	25	342	0.85460888	9.511321127	3.476309529	Up	9.32E-12	2.08E-11	AAEL000287-PA (Fragment) OS=Aedes aegypti GN=AAEL000287 PE=4 SV=1
CCG010791.1	759	42	342	2.37777188	15.7519508	2.727848296	Up	7.66E-14	2.47E-13	AAEL006747-PA OS=Aedes aegypti GN=AAEL006747 PE=4 SV=1
CCG001113.1	759	54	342	3.05713527	15.7519508	2.365278217	Up	0	0	AAEL000865-PA OS=Aedes aegypti GN=AAEL000865 PE=4 SV=1
CCG012685.1	906	74	342	3.50966927	13.1961707	1.910712353	Up	0	0	AAEL002179-PA OS=Aedes aegypti GN=AAEL002179 PE=4 SV=1
CCG005766.1	693	28	343	1.73615089	17.30258143	3.317023048	Up	1.17E-12	2.86E-12	AAEL007053-PB OS=Aedes aegypti GN=AAEL007053 PE=4 SV=1
CCG016809.1	1788	49	343	1.17758221	6.706201864	2.509668126	Up	0	0	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010644 PE=4 SV=1
CCG010413.1	1905	54	343	1.21803972	6.294324899	2.369490468	Up	0	0	AAEL004917-PA OS=Aedes aegypti GN=AAEL004917 PE=4 SV=1
CCG010380.1	561	107	343	8.19565348	21.37377706	1.382910984	Up	0	0	AAEL009850-PA OS=Aedes aegypti GN=GALE14 PE=4 SV=1

CCG018692.1	2663	123	343	1.98470798	4.502699562	1.181863465	Up	0	0	AAEL003974-PA OS=Aedes aegypti GN=AAEL003974 PE=4 SV=1
CCG018943.2	1554	607	343	16.7841885	7.716016044	-1.121174736	Down	4.78E-32	2.03E-31	Cytochrome P450 CYP6N3v2 OS=Aedes albopictus PE=2 SV=1
CCG003080.1	420	14	344	1.43232449	28.63249336	4.321223036	Up	3.07E-08	6.01E-08	-
CCG023734.1	829	71	344	3.68015821	14.50620894	1.978830839	Up	0	0	Alkaline phosphatase OS=Aedes aegypti GN=AAEL003298 PE=3 SV=1
CCG001483.1	966	105	344	4.67062333	12.44891015	1.414332441	Up	0	0	Conserved membrane protein OS=Aedes aegypti PE=2 SV=1
CCG027252.1	1260	116	344	3.95594382	9.544164452	1.270596963	Up	1.16E-13	3.58E-13	AAEL013825-PA OS=Aedes aegypti GN=AAEL013825 PE=4 SV=1
CCG028518.1	708	744	344	45.1546364	16.98537741	-1.410580853	Down	4.06E-55	1.93E-54	AAEL006131-PA OS=Aedes aegypti GN=AAEL006131 PE=4 SV=1
CCG019278.1	777	5	345	0.27651052	15.52201478	5.810837661	Up	1.59E-05	2.76E-05	AAEL011383-PA OS=Aedes aegypti GN=AAEL011383 PE=4 SV=1
CCG015233.1	1914	70	345	1.57151589	6.301256785	2.003482739	Up	0	0	AAEL009262-PA OS=Aedes aegypti GN=AAEL009262 PE=4 SV=1
CCG002374.1	669	72	345	4.62454543	18.02781089	1.962840754	Up	0	0	AAEL014052-PA OS=Aedes aegypti GN=AAEL014052 PE=4 SV=1
CCG012530.4	2316	9883	345	183.363509	5.207515322	-5.13796757	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008688 PE=4 SV=1
CCG016272.1	2004	5	346	0.10720992	6.035710461	5.815013337	Up	1.59E-05	2.76E-05	AAEL004909-PA OS=Aedes aegypti GN=AAEL004909 PE=4 SV=1
CCG026622.1	1071	12	346	0.48145361	11.29371033	4.551978931	Up	1.30E-07	2.49E-07	AAEL004943-PA OS=Aedes aegypti GN=AAEL004943 PE=4 SV=1
CCG020339.1	1254	23	346	0.78812113	9.645585138	3.613379475	Up	4.01E-11	8.67E-11	AAEL014744-PA OS=Aedes aegypti GN=AAEL014744 PE=4 SV=1
CCG002422.1	1896	28	346	0.63457414	6.379516753	3.329586509	Up	1.17E-12	2.85E-12	AAEL009031-PA OS=Aedes aegypti GN=AAEL009031 PE=4 SV=1
CCG012945.1	1829	47	346	1.10419766	6.613211462	2.58235258	Up	0	0	AAEL008643-PA OS=Aedes aegypti GN=AAEL008643 PE=4 SV=1

CCG023126.1	1467	54	346	1.58170802	8.245101406	2.382053929	Up	0	0	Serine/threonine-protein kinase RIO1 OS=Aedes aegypti GN=AAEL011857 PE=3 SV=1
CCG014620.1	912	65	346	3.06253591	13.26267956	2.114573618	Up	0	0	AAEL005840-PA OS=Aedes aegypti GN=AAEL005846 PE=4 SV=1
CCG016866.1	3342	73	346	0.93859684	3.619259055	1.947116873	Up	0	0	AAEL009289-PA OS=Aedes aegypti GN=AAEL009289 PE=4 SV=1
CCG000308.1	5385	105	346	0.83784998	2.246158545	1.422695914	Up	0	0	AAEL012462-PA OS=Aedes aegypti GN=AAEL012462 PE=4 SV=1
CCG016306.1	1018	126	346	5.31845438	11.88169328	1.159661508	Up	1.34E-13	4.06E-13	AAEL006010-PA OS=Aedes aegypti GN=AAEL006010 PE=4 SV=1
CCG010798.1	659	140	346	9.12862344	18.35442149	1.007658415	Up	4.03E-13	1.07E-12	AAEL007825-PA OS=Aedes aegypti GN=AAEL007825 PE=4 SV=1
CCG013959.1	675	714	346	45.4524304	17.91935372	-1.342838833	Down	3.12E-49	1.45E-48	AAEL001946-PA OS=Aedes aegypti GN=AAEL001946 PE=4 SV=1
CCG009996.1	1476	2573	346	74.9059127	8.194826398	-3.19229431	Down	0	0	AAEL004440-PA OS=Aedes aegypti GN=AAEL004440 PE=4 SV=1
CCG016497.1	410	7	347	0.73362962	29.58663912	5.333750134	Up	4.31E-06	7.71E-06	AAEL005153-PA OS=Aedes aegypti GN=AAEL005153 PE=4 SV=1
CCG009342.1	1011	14	347	0.59503094	11.99853812	4.333750134	Up	3.07E-08	6.00E-08	AAEL003107-PA OS=Aedes aegypti GN=AAEL003107 PE=4 SV=1
CCG019473.1	902	25	347	1.19095717	13.44847233	3.497248867	Up	9.32E-12	2.08E-11	AAEL012372-PA OS=Aedes aegypti GN=AAEL012372 PE=4 SV=1
CCG006456.1	1107	39	347	1.51383889	10.95801449	2.855702838	Up	7.33E-14	2.38E-13	AAEL000278-PA (Fragment) OS=Aedes aegypti GN=AAEL000278 PE=3 SV=1
CCG027503.1	1269	53	347	1.79463825	9.559119023	2.413184602	Up	0	0	AAEL010770-PA OS=Aedes aegypti GN=AAEL010770 PE=4 SV=1
CCG020473.1	1199	80	347	2.86703817	10.11719937	1.819176962	Up	0	0	AAEL001407-PA OS=Aedes aegypti GN=AAEL001407 PE=4 SV=1
CCG025841.1	705	109	347	6.64354762	17.20641424	1.372920732	Up	0	0	Hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG008384.1	2436	119	347	2.09909623	4.979688851	1.246287293	Up	0	0	AAEL005402-PA OS=Aedes aegypti GN=AAEL005402 PE=4 SV=1

CCG001821.1	787	17	348	0.92818995	15.4580436	4.057793858	Up	3.39E-09	6.89E-09	AAEL002905-PA OS=Aedes aegypti GN=AAEL002905 PE=4 SV=1
CCG004516.1	780	31	348	1.7077715	15.59676964	3.191060389	Up	2.40E-13	6.76E-13	AAEL001684-PA OS=Aedes aegypti GN=AAEL001684 PE=4 SV=1
CCG026989.1	3828	57	348	0.63983147	3.178025161	2.312366686	Up	0	0	AAEL010355-PA OS=Aedes aegypti GN=AAEL010355 PE=4 SV=1
CCG003112.1	905	76	348	3.6085081	13.44251969	1.897329186	Up	0	0	AAEL008755-PA OS=Aedes aegypti GN=AAEL008755 PE=4 SV=1
CCG017680.1	1833	78	348	1.82849935	6.63692325	1.859854481	Up	0	0	AAEL006640-PA OS=Aedes aegypti GN=AAEL006640 PE=3 SV=1
CCG011839.1	1698	106	348	2.68244515	7.164593826	1.417336245	Up	0	0	AAEL004868-PA OS=Aedes aegypti GN=AAEL004868 PE=4 SV=1
CCG010957.1	2115	126	348	2.55989908	5.75200015	1.167976776	Up	1.34E-13	4.06E-13	AGAP009971-PA (Fragment) OS=Anopheles gambiae GN=AGAP009971 PE=3 SV=3
CCG007577.1	2527	131	348	2.22755648	4.81419878	1.111833698	Up	1.29E-13	3.96E-13	AAEL007729-PA OS=Aedes aegypti GN=AAEL007729 PE=4 SV=1
CCG013708.1	1566	605	348	16.6006957	7.768505949	-1.095534632	Down	7.91E-31	3.35E-30	AAEL000714-PA OS=Aedes aegypti GN=AAEL000714 PE=4 SV=1
CCG012327.1	923	819	348	38.1280744	13.18036871	-1.532462942	Down	2.64E-68	1.30E-67	AAEL000692-PA OS=Aedes aegypti GN=AAEL000692 PE=4 SV=1
CCG019888.1	2049	1596	348	33.4698372	5.937276875	-2.494988237	Down	4.03E-251	2.48E-250	AAEL014457-PA (Fragment) OS=Aedes aegypti GN=AAEL014457 PE=4 SV=1
CCG014681.1	1476	28	349	0.81514402	8.265879806	3.342041508	Up	1.17E-12	2.85E-12	AAEL000699-PA OS=Aedes aegypti GN=AAEL000699 PE=4 SV=1
CCG005346.1	1917	41	349	0.91901884	6.364339381	2.791844425	Up	1.02E-13	3.19E-13	AAEL010136-PA OS=Aedes aegypti GN=AAEL010136 PE=4 SV=1
CCG012270.2	691	48	349	2.98487303	17.65620636	2.564433929	Up	5.35E-14	1.79E-13	AAEL010182-PA OS=Aedes aegypti GN=AAEL010182 PE=4 SV=1
CCG014733.1	834	876	349	45.1336781	14.62882325	-1.62539063	Down	2.27E-79	1.15E-78	AAEL001031-PA OS=Aedes aegypti GN=AAEL001031 PE=3 SV=1
CCG025057.1	714	83	350	4.99508119	17.13641018	1.778484884	Up	1.32E-13	4.02E-13	Dacapo OS=Aedes albopictus PE=2 SV=1

CCG027685.1	591	8635	350	627.823449	20.70287118	-4.922456147	Down	0	0	AAEL009707-PA OS=Aedes aegypti GN=AAEL009707 PE=4 SV=1
CCG025257.1	1843	40	351	0.93260412	6.657816141	2.835712329	Up	6.86E-14	2.25E-13	AAEL007461-PA OS=Aedes aegypti GN=AAEL007461 PE=4 SV=1
CCG013852.1	810	85	351	4.50916968	15.1485866	1.748249488	Up	0	0	AAEL001664-PA OS=Aedes aegypti GN=AAEL001664 PE=4 SV=1
CCG006132.1	2694	919	351	14.6581983	4.55469753	-1.686280627	Down	1.27E-87	6.59E-87	AAEL003472-PA OS=Aedes aegypti GN=AAEL003472 PE=4 SV=1
CCG009528.1	2076	1559	351	32.2686976	5.910575697	-2.448764789	Down	5.30E-240	3.25E-239	AAEL011436-PA OS=Aedes aegypti GN=AAEL011436 PE=4 SV=1
CCG014821.1	348	11	352	1.35823874	35.36009605	4.702313204	Up	2.67E-07	5.03E-07	AAEL007291-PA OS=Aedes aegypti GN=AAEL007291 PE=4 SV=1
CCG004295.1	1476	49	352	1.42650203	8.336933214	2.547034978	Up	0	0	AAEL001359-PA OS=Aedes aegypti GN=AAEL001359 PE=4 SV=1
CCG009428.1	2319	86	352	1.59353048	5.306301606	1.735480068	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007254 PE=4 SV=1
CCG017981.1	969	2038	352	90.3739104	12.69898186	-2.831193514	Down	0	0	AAEL004118-PA OS=Aedes aegypti GN=AAEL004118 PE=4 SV=1
CCG010905.1	442	0	353	0.001	27.91916674	14.76896826	Up	4.44E-16	1.64E-15	Predicted protein OS=Nematostella vectensis GN=v1g229551 PE=3 SV=1
CCG027194.1	1626	53	353	1.40061251	7.589342989	2.437917123	Up	0	0	AAEL003657-PA OS=Aedes aegypti GN=AAEL003657 PE=4 SV=1
CCG022161.1	2090	60	353	1.2335809	5.90443622	2.258946981	Up	0	0	AAEL013550-PA OS=Aedes aegypti GN=AAEL013550 PE=4 SV=1
CCG026192.1	1464	67	353	1.96651108	8.429147336	2.099748387	Up	0	0	AAEL014112-PA OS=Aedes aegypti GN=AAEL014112 PE=4 SV=1
CCG003173.1	618	79	353	5.49289488	19.96807719	1.862056829	Up	0	0	AAEL002898-PA OS=Aedes aegypti GN=AAEL002898 PE=3 SV=1
CCG011833.1	2184	79	353	1.55430817	5.650307555	1.862056829	Up	0	0	AAEL004889-PA OS=Aedes aegypti GN=AAEL004889 PE=4 SV=1
CCG001456.1	623	79	353	5.44881065	19.80781974	1.862056829	Up	0	0	AAEL008355-PA OS=Aedes aegypti GN=AAEL008355 PE=4 SV=1

CCG009425.1	927	91	353	4.21817244	13.31205146	1.658042937	Up	0	0	AAEL008120-PA OS=Aedes aegypti GN=AAEL008120 PE=3 SV=1
CCG020686.1	927	130	353	6.02596063	13.31205146	1.143469764	Up	0	0	AGAP008088-PA OS=Anopheles gambiae GN=AGAP008088 PE=4 SV=4
CCG010137.1	477	0	354	0.001	25.94387836	14.66310654	Up	4.44E-16	1.64E-15	Histone H2B OS=Aedes aegypti GN=AAEL015674 PE=3 SV=1
CCG002165.1	960	43	354	1.92468603	12.89086456	2.743653999	Up	8.46E-14	2.70E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013990 PE=4 SV=1
CCG011281.1	873	47	354	2.31337632	14.17552117	2.615329902	Up	0	0	AAEL003342-PA OS=Aedes aegypti GN=AAEL003342 PE=4 SV=1
CCG009068.1	2175	85	354	1.67927698	5.689760909	1.760527818	Up	0	0	AAEL007630-PA OS=Aedes aegypti GN=AAEL007630 PE=4 SV=1
CCG009327.1	1344	130	354	4.15629873	9.2077604	1.147550941	Up	0	0	AAEL003964-PA OS=Aedes aegypti GN=AAEL003964 PE=4 SV=1
CCG023009.1	3257	661	354	8.72060012	3.799579361	-1.198587708	Down	1.68E-38	7.43E-38	AAEL001431-PA OS=Aedes aegypti GN=AAEL001431 PE=4 SV=1
CCG023637.1	732	33	355	1.93716017	16.95380909	3.129594299	Up	1.31E-13	4.02E-13	Serehiac OS=Culex quinquefasciatus GN=CpipJ_CPIJ001564 PE=4 SV=1
CCG002237.1	1830	42	355	0.98619063	6.781523636	2.781670995	Up	7.66E-14	2.47E-13	AAEL009306-PA OS=Aedes aegypti GN=AAEL009306 PE=4 SV=1
CCG016860.1	1956	49	355	1.07644018	6.344677022	2.559278574	Up	0	0	AAEL003660-PA OS=Aedes aegypti GN=AAEL003660 PE=4 SV=1
CCG025983.1	2013	68	355	1.45153599	6.165021487	2.086525577	Up	0	0	AAEL002013-PA OS=Aedes aegypti GN=AAEL002013 PE=4 SV=1
CCG023798.1	918	142	355	6.64673455	13.51872359	1.024241299	Up	3.60E-13	9.69E-13	AAEL015041-PA OS=Aedes aegypti GN=AAEL015041 PE=3 SV=1
CCG021652.1	831	699	355	36.1442172	14.93404122	-1.275160227	Down	1.25E-44	5.71E-44	-
CCG010019.1	7101	2913	355	17.6272126	1.747667688	-3.334301568	Down	0	0	AAEL012268-PA OS=Aedes aegypti GN=AAEL012268 PE=4 SV=1
CCG006626.1	972	45	356	1.98933957	12.80364869	2.686193538	Up	0	0	AAEL009399-PA OS=Aedes aegypti GN=AAEL009399 PE=4 SV=1

CCG026158.1	939	47	356	2.15077479	13.25361718	2.623457783	Up	0	0	AAEL003839-PA OS=Aedes aegypti GN=AAEL003839 PE=4 SV=1
CCG025899.1	846	52	356	2.64116572	14.7105751	2.477606917	Up	0	0	AAEL011069-PA OS=Aedes aegypti GN=AAEL011069 PE=4 SV=1
CCG004184.1	913	99	356	4.65936881	13.63104768	1.548690015	Up	0	0	AAEL012525-PA OS=Aedes aegypti GN=AAEL012525 PE=4 SV=1
CCG012611.1	1394	121	356	3.72979762	8.927651744	1.259183398	Up	0	0	AAEL007574-PA OS=Aedes aegypti GN=AAEL007574 PE=4 SV=1
CCG017920.1	2883	598	356	8.91290368	4.316734836	-1.045955039	Down	2.21E-28	9.24E-28	AAEL001225-PA OS=Aedes aegypti GN=AAEL001225 PE=3 SV=1
CCG026700.2	1210	703	356	24.9650607	10.28524507	-1.279334244	Down	4.27E-45	1.95E-44	AAEL013984-PA OS=Aedes aegypti GN=AAEL013984 PE=4 SV=1
CCG018505.1	1143	2	357	0.07518764	10.91872687	7.182093468	Up	9.36E-05	0.00015385	AAEL005938-PA OS=Aedes aegypti GN=AAEL005938 PE=4 SV=1
CCG004545.1	744	38	357	2.19469075	16.77433442	2.934165954	Up	4.26E-14	1.45E-13	-
CCG026488.1	1953	71	357	1.56213577	6.390222636	2.032346348	Up	0	0	AAEL004950-PA OS=Aedes aegypti GN=AAEL004950 PE=4 SV=1
CCG007949.1	1513	72	357	2.04482544	8.24858216	2.012168466	Up	0	0	AAEL003101-PA OS=Aedes aegypti GN=AAEL003101 PE=4 SV=1
CCG010460.1	1272	73	357	2.46603037	9.811403151	1.992268909	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_22896 PE=4 SV=1
CCG005847.1	1179	142	357	5.17532003	10.58533063	1.032346348	Up	3.20E-13	8.78E-13	AAEL003776-PA OS=Aedes aegypti GN=AAEL003776 PE=4 SV=1
CCG002287.1	2724	4	358	0.063098	4.594369708	6.186128981	Up	2.95E-05	5.04E-05	AAEL004462-PA OS=Aedes aegypti GN=AAEL004462 PE=4 SV=1
CCG012073.1	2616	55	358	0.90341567	4.784045522	2.404769268	Up	0	0	AAEL009438-PA OS=Aedes aegypti GN=AAEL009438 PE=4 SV=1
CCG010284.1	4336	106	358	1.05045938	2.886315287	1.458208527	Up	0	0	AAEL014493-PA OS=Aedes aegypti GN=AAEL014493 PE=4 SV=1
CCG000635.2	6188	115	358	0.79856488	2.022473026	1.34063893	Up	6.13E-14	2.03E-13	AAEL005723-PA OS=Aedes aegypti GN=AAEL005723 PE=4 SV=1
CCG022209.1	2254	139	358	2.64986385	5.552379363	1.067187908	Up	1.57E-13	4.66E-13	AAEL000899-PA OS=Aedes aegypti GN=AAEL000899 PE=4 SV=1

CCG006802.1	4467	663	358	6.37764362	2.801670715	-1.186736079	Down	4.96E-38	2.19E-37	AAEL012249-PA (Fragment) OS=Aedes aegypti GN=AAEL012249 PE=4 SV=1
CCG010818.1	933	29	359	1.33560804	13.45125548	3.332172243	Up	6.66E-13	1.69E-12	Putative salivary Gram negative bacteria-binding protein OS=Aedes albopictus PE=2 SV=1
CCG013420.1	1785	38	359	0.91476186	7.030824292	2.942225724	Up	4.26E-14	1.45E-13	AAEL004654-PA OS=Aedes aegypti GN=AAEL004654 PE=4 SV=1
CCG015133.1	2100	58	359	1.18678315	5.976200648	2.332172243	Up	0	0	AAEL011415-PA OS=Aedes aegypti GN=AAEL011415 PE=4 SV=1
CCG008484.1	1176	92	359	3.36157788	10.67178687	1.666591282	Up	1.44E-14	5.12E-14	Putative salivary Gram negative bacteria-binding protein OS=Aedes albopictus PE=2 SV=1
CCG023146.1	699	127	359	7.80709055	17.95425087	1.201468551	Up	1.57E-13	4.66E-13	AAEL008007-PA OS=Aedes aegypti GN=AAEL008007 PE=4 SV=1
CCG016999.1	942	137	359	6.24931384	13.3227403	1.092121155	Up	2.92E-13	8.09E-13	AAEL006853-PA OS=Aedes aegypti GN=AAEL006853 PE=4 SV=1
CCG028019.1	753	140	359	7.98906088	16.66669503	1.060870221	Up	8.88E-16	3.24E-15	AAEL013188-PA OS=Aedes aegypti GN=AAEL013188 PE=4 SV=1
CCG012139.3	1176	2048	359	74.8316467	10.67178687	-2.809846762	Down	0	0	Takeout OS=Aedes aegypti PE=2 SV=1
CCG026694.1	681	48	360	3.02870376	18.48014631	2.609203799	Up	5.35E-14	1.80E-13	AAEL004438-PA OS=Aedes aegypti GN=AAEL004438 PE=3 SV=1
CCG002139.1	1212	77	360	2.72992538	10.38364657	1.927379759	Up	0	0	AAEL010427-PA OS=Aedes aegypti GN=AAEL010427 PE=4 SV=1
CCG002792.1	6775	96	360	0.60887004	1.85756157	1.609203799	Up	0	0	AGAP001529-PA OS=Anopheles gambiae GN=AgaP_AGAP001529 PE=4 SV=5
CCG012981.1	1299	940	360	31.0943422	9.688206034	-1.682350646	Down	2.75E-89	1.43E-88	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009026 PE=4 SV=1
CCG008204.1	1221	90	361	3.16730231	10.33573949	1.706315134	Up	0	0	AAEL012942-PA OS=Aedes aegypti GN=AAEL012942 PE=4 SV=1

CCG013462.1	1817	735	361	17.3818134	6.945480415	-1.323432209	Down	1.55E-49	7.23E-49	AAEL006305-PA (Fragment) OS=Aedes aegypti GN=AAEL006305 PE=4 SV=1
CCG003415.1	1650	1263	361	32.8913787	7.648447221	-2.104470693	Down	1.50E-161	8.67E-161	AAEL008635-PA OS=Aedes aegypti GN=AAEL008635 PE=3 SV=1
CCG028204.1	3018	1616	361	23.0083138	4.181556632	-2.460043252	Down	4.65E-250	2.86E-249	AAEL014426-PA OS=Aedes aegypti GN=AAEL014426 PE=3 SV=1
CCG027198.1	489	0	362	0.001	25.87913332	14.65950168	Up	4.44E-16	1.64E-15	Predicted protein OS=Nematostella vectensis GN=v1g229551 PE=3 SV=1
CCG022985.1	1215	39	362	1.37927543	10.41555242	2.916756872	Up	7.33E-14	2.38E-13	AAEL014267-PA OS=Aedes aegypti GN=AAEL014267 PE=4 SV=1
CCG008180.1	594	43	362	3.11060368	21.30453904	2.775894336	Up	8.46E-14	2.70E-13	AAEL006290-PA OS=Aedes aegypti GN=AAEL006290 PE=4 SV=1
CCG000812.1	1749	59	362	1.44952221	7.235503826	2.319516042	Up	0	0	AAEL015533-PA OS=Aedes aegypti GN=AAEL015533 PE=3 SV=1
CCG003274.1	2468	78	362	1.35803861	5.12759165	1.916756872	Up	0	0	AI_G2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ015325 PE=3 SV=1
CCG003036.1	465	114	362	10.5345156	27.21483052	1.369269077	Up	8.48E-14	2.70E-13	AAEL013494-PA OS=Aedes aegypti GN=AAEL013494 PE=4 SV=1
CCG020562.1	848	114	362	5.77659168	14.92322664	1.369269077	Up	8.48E-14	2.70E-13	AAEL007992-PA OS=Aedes aegypti GN=AAEL007992 PE=3 SV=1
CCG019801.1	2352	146	362	2.66733897	5.380483075	1.012334532	Up	1.05E-13	3.29E-13	AAEL007753-PA OS=Aedes aegypti GN=AAEL007753 PE=4 SV=1
CCG020112.1	864	17	363	0.84546932	14.68733156	4.118676101	Up	3.39E-09	6.90E-09	-
CCG006610.1	1369	21	363	0.65914129	9.269433505	3.813821519	Up	1.76E-10	3.72E-10	AAEL014934-PA OS=Aedes aegypti GN=AAEL014934 PE=4 SV=1
CCG011205.1	539	67	363	5.34132137	23.54332925	2.140049751	Up	0	0	AAEL013291-PA OS=Aedes aegypti GN=AAEL013291 PE=4 SV=1
CCG028002.1	776	940	363	52.0509672	16.35290524	-1.670378005	Down	2.16E-88	1.12E-87	AAEL007489-PA OS=Aedes aegypti GN=AAEL007489 PE=4 SV=1
CCG025602.1	783	1529	363	83.9089709	16.20671069	-2.37223375	Down	1.19E-226	7.25E-226	AAEL004957-PA OS=Aedes aegypti GN=AAEL004957 PE=4 SV=1
CCG011436.1	490	2184	363	191.522246	25.89766218	-2.886618199	Down	0	0	AAEL000958-PA OS=Aedes aegypti GN=AAEL000958 PE=4 SV=1

CCG017801.1	1556	27	364	0.74561879	8.177900222	3.455220342	Up	2.28E-12	5.41E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017880 PE=4 SV=1
CCG020068.1	1110	74	364	2.86464897	11.46379527	2.000654478	Up	0	0	AAEL010605-PA (Fragment) OS=Aedes aegypti GN=AAEL010605 PE=4 SV=1
CCG015410.1	464	1298	364	120.204128	27.4241654	-2.131966824	Down	9.29E-169	5.38E-168	AAEL004060-PA OS=Aedes aegypti GN=AAEL004060 PE=4 SV=1
CCG018133.1	1971	1656	364	36.1024254	6.456018643	-2.483379113	Down	4.59E-259	2.84E-258	AAEL014156-PA OS=Aedes aegypti GN=AAEL014156 PE=4 SV=1
CCG018335.1	1437	30	365	0.8970717	8.879450955	3.307175262	Up	3.71E-13	9.95E-13	AAEL007774-PA OS=Aedes aegypti GN=AAEL007774 PE=4 SV=1
CCG001698.1	1029	34	365	1.41979687	12.4001662	3.126603016	Up	8.82E-14	2.79E-13	AAEL007279-PA OS=Aedes aegypti GN=AAEL007279 PE=3 SV=1
CCG009024.1	612	66	365	4.63399099	20.84929906	2.169671738	Up	0	0	AAEL007211-PA OS=Aedes aegypti GN=AAEL007211 PE=4 SV=1
CCG026380.1	1153	75	365	2.79508248	11.06658371	1.985247167	Up	0	0	AAEL009465-PA OS=Aedes aegypti GN=AAEL009465 PE=4 SV=1
CCG024325.1	1914	107	365	2.4021743	6.666547033	1.472598871	Up	0	0	AAEL008161-PA (Fragment) OS=Aedes aegypti GN=AAEL008161 PE=4 SV=1
CCG023933.1	648	109	365	7.22793375	19.69100466	1.445881533	Up	0	0	AAEL010046-PA OS=Aedes aegypti GN=AAEL010046 PE=4 SV=1
CCG018063.1	1124	133	365	5.08449707	11.35210945	1.158783422	Up	9.90E-14	3.11E-13	AAEL001597-PA OS=Aedes aegypti GN=AAEL001597 PE=4 SV=1
CCG022503.1	1455	135	365	3.98688259	8.769602077	1.137250261	Up	1.90E-13	5.49E-13	AAEL002285-PA (Fragment) OS=Aedes aegypti GN=AAEL002285 PE=4 SV=1
CCG006922.1	852	1027	365	51.7956778	14.97625707	-1.790154609	Down	3.44E-106	1.84E-105	AAEL014354-PA OS=Aedes aegypti GN=CLIPB43 PE=3 SV=1
CCG007182.1	915	30	366	1.40884376	13.98331071	3.311122446	Up	3.71E-13	9.93E-13	-
CCG021636.1	834	45	366	2.31851086	15.34140204	2.726159946	Up	0	0	Ecdysone-induced protein 75C OS=Aedes aegypti GN=e75C PE=2 SV=1

CCG008762.1	978	87	366	3.82246105	13.0825453	1.775069546	Up	0	0	Nuclear distribution protein NUDC OS=Aedes aegypti PE=2 SV=1
CCG021942.1	1170	112	366	4.11334212	10.93566607	1.41065812	Up	0	0	AGAP004234-PA OS=Anopheles gambiae GN=AgaP_AGAP004234 PE=4 SV=4
CCG019457.1	5001	140	366	1.20291199	2.558434173	1.088730025	Up	0	0	AAEL009806-PA (Fragment) OS=Aedes aegypti GN=AAEL009806 PE=4 SV=1
CCG006702.1	1740	769	366	18.9906471	7.3532927	-1.368826746	Down	2.00E-54	9.47E-54	AAEL000993-PA OS=Aedes aegypti GN=AAEL000993 PE=3 SV=1
CCG001899.1	4206	792	366	8.09130523	3.042018378	-1.411343578	Down	1.32E-58	6.33E-58	AAEL010379-PA OS=Aedes aegypti GN=AAEL010379 PE=3 SV=1
CCG028309.1	727	892	366	52.7221503	17.59935254	-1.582886858	Down	8.78E-78	4.44E-77	-
CCG023564.1	614	2	367	0.13996656	20.89525664	7.221949457	Up	9.36E-05	0.00015322	GK23465 OS=Drosophila willistoni GN=GK23465 PE=4 SV=1
CCG006818.1	246	21	367	3.66814808	52.15320153	3.829632034	Up	1.76E-10	3.72E-10	AAEL007539-PA OS=Aedes aegypti GN=AAEL007539 PE=4 SV=1
CCG016284.1	767	27	367	1.5126243	16.72710245	3.467061954	Up	2.28E-12	5.41E-12	AAEL000697-PA OS=Aedes aegypti GN=AAEL000697 PE=4 SV=1
CCG027995.1	1962	42	367	0.91984141	6.53908643	2.829632034	Up	7.66E-14	2.47E-13	AAEL000685-PA OS=Aedes aegypti GN=AAEL000685 PE=4 SV=1
CCG021347.1	926	62	367	2.87702327	13.85495419	2.267753146	Up	0	0	AAEL003360-PA OS=Aedes aegypti GN=AAEL003360 PE=4 SV=1
CCG010026.1	1016	64	367	2.70675494	12.62764525	2.221949457	Up	0	0	AAEL003390-PA OS=Aedes aegypti GN=AAEL003390 PE=4 SV=1
CCG018164.1	1383	69	367	2.14382624	9.276708297	2.113425	Up	0	0	AAEL002427-PA OS=Aedes aegypti GN=AAEL002427 PE=4 SV=1
CCG020621.1	1788	999	367	24.0082578	7.175440478	-1.742391411	Down	1.51E-99	8.01E-99	AGAP002896-PA (Fragment) OS=Anopheles gambiae GN=AgaP_AGAP002896 PE=4 SV=5
CCG004423.1	480	3	368	0.26856084	26.80134553	6.640912659	Up	5.28E-05	8.84E-05	Putative salivary protein 34k-1 OS=Aedes albopictus PE=2 SV=1
CCG027390.1	1636	41	368	1.07686988	7.86347546	2.868323155	Up	1.02E-13	3.19E-13	AAEL002852-PA OS=Aedes aegypti GN=AAEL002852 PE=4 SV=1

CCG014346.1	1866	41	368	0.94413672	6.894236791	2.868323155	Up	1.02E-13	3.19E-13	AAEL014291-PA OS=Aedes aegypti GN=AAEL014291 PE=4 SV=1
CCG024130.1	924	50	368	2.32520209	13.9227769	2.58201897	Up	0	0	AAEL004519-PA OS=Aedes aegypti GN=AAEL004519 PE=4 SV=1
CCG011144.1	5088	81	368	0.68407007	2.528428823	1.886025157	Up	4.46E-14	1.51E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016906 PE=4 SV=1
CCG028394.1	1145	111	368	4.16562493	11.23549856	1.431459294	Up	0	0	AAEL000702-PA OS=Aedes aegypti GN=AAEL000702 PE=4 SV=1
CCG013044.1	1440	128	368	3.81953197	8.933781842	1.22587516	Up	4.15E-14	1.42E-13	UDP-glucose 6-dehydrogenase OS=Aedes aegypti GN=AAEL011242 PE=3 SV=1
CCG005634.1	2049	658	368	13.798968	6.278499684	-1.136068614	Down	2.61E-35	1.13E-34	AAEL007030-PA OS=Aedes aegypti GN=AAEL007030 PE=4 SV=1
CCG025112.1	825	2634	368	137.190644	15.59351012	-3.13716447	Down	0	0	AAEL003474-PA OS=Aedes aegypti GN=AAEL003474 PE=4 SV=1
CCG019723.1	836	25	369	1.2849801	15.43014848	3.58593402	Up	9.32E-12	2.08E-11	AAEL013351-PA OS=Aedes aegypti GN=AAEL013351 PE=3 SV=1
CCG000291.1	1379	40	369	1.24640274	9.354317715	2.907862115	Up	6.86E-14	2.24E-13	-
CCG010218.1	1438	56	369	1.67336936	8.970517475	2.422435288	Up	0	0	AAEL010331-PA OS=Aedes aegypti GN=AAEL010331 PE=4 SV=1
CCG022089.1	853	77	369	3.87886233	15.12263087	1.963003669	Up	0	0	AAEL004291-PA OS=Aedes aegypti GN=AAEL004291 PE=4 SV=1
CCG003536.1	1398	77	369	2.36671643	9.227184642	1.963003669	Up	0	0	AAEL002044-PA OS=Aedes aegypti GN=AAEL002044 PE=4 SV=1
CCG009861.1	501	97	369	8.31948954	25.74771283	1.629877368	Up	3.62E-14	1.25E-13	AAEL014545-PA OS=Aedes aegypti GN=AAEL014545 PE=4 SV=1
CCG025880.1	884	117	369	5.68717076	14.59231236	1.35942549	Up	9.44E-14	2.98E-13	AAEL009284-PA OS=Aedes aegypti GN=AAEL009284 PE=4 SV=1
CCG013002.3	1044	136	369	5.59758995	12.35594265	1.142327369	Up	1.70E-13	4.99E-13	AAEL013065-PA OS=Aedes aegypti GN=AAEL013065 PE=4 SV=1
CCG010470.1	903	611	369	29.0747595	14.28527589	-1.02523836	Down	4.49E-28	1.87E-27	-
CCG004611.1	4053	676	369	7.16692341	3.182729862	-1.171089226	Down	5.80E-38	2.56E-37	AAEL006522-PA (Fragment) OS=Aedes aegypti GN=AAEL006522 PE=4 SV=1

CCG008545.1	522	12	371	0.98780999	24.84582506	4.65262608	Up	1.30E-07	2.49E-07	-
CCG003818.1	1116	28	371	1.0780937	11.6214343	3.430233658	Up	1.17E-12	2.85E-12	AAEL004955-PA OS=Aedes aegypti GN=AAEL004955 PE=4 SV=1
CCG006919.1	2324	36	371	0.66562412	5.580688762	3.067663579	Up	6.13E-14	2.03E-13	DNA ligase (Fragment) OS=Anopheles gambiae GN=AGAP009222 PE=3 SV=3
CCG007214.1	965	37	371	1.64754423	13.43991781	3.028135215	Up	5.17E-14	1.74E-13	-
CCG008338.2	1283	87	371	2.91377	10.10874566	1.794645085	Up	0	0	AAEL004309-PA OS=Aedes aegypti GN=AAEL004309 PE=4 SV=1
CCG008800.1	1206	88	371	3.13543669	10.75416309	1.778156962	Up	2.93E-14	1.02E-13	AAEL009170-PA OS=Aedes aegypti GN=AAEL009170 PE=3 SV=1
CCG002518.1	873	129	371	6.34947968	14.85626653	1.226361325	Up	0	0	AAEL005479-PA OS=Aedes aegypti GN=AAEL005479 PE=4 SV=1
CCG001200.1	1944	1083	371	23.9383861	6.671564137	-1.843228947	Down	1.89E-116	1.03E-115	AAEL005160-PA OS=Aedes aegypti GN=AAEL005160 PE=4 SV=1
CCG024809.1	1590	18	372	0.48644983	8.178917584	4.071547013	Up	1.62E-09	3.32E-09	AGAP004522-PA OS=Anopheles gambiae GN=AgaP_AGAP004522 PE=4 SV=4
CCG003887.1	1809	31	372	0.73635256	7.188766699	3.287275705	Up	2.40E-13	6.77E-13	AAEL013167-PA OS=Aedes aegypti GN=AAEL013167 PE=4 SV=1
CCG021040.1	1182	60	372	2.1812048	11.00209726	2.334581419	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008082 PE=4 SV=1
CCG003001.1	1053	124	372	5.06006371	12.34993253	1.287275705	Up	1.61E-13	4.76E-13	AAEL015291-PA (Fragment) OS=Aedes aegypti GN=AAEL015291 PE=4 SV=1
CCG026512.1	877	629	372	30.818658	14.82836825	-1.055444192	Down	3.45E-30	1.46E-29	AAEL010002-PA OS=Aedes aegypti GN=AAEL010002 PE=4 SV=1
CCG000018.1	3901	2428	372	26.7445567	3.333627008	-3.004080691	Down	0	0	AAEL004133-PA OS=Aedes aegypti GN=AAEL004133 PE=4 SV=1
CCG018508.1	1461	23	373	0.67645715	8.925008375	3.721783068	Up	4.01E-11	8.66E-11	AAEL001870-PA OS=Aedes aegypti GN=AAEL001870 PE=4 SV=1
CCG001396.1	507	48	373	4.06814055	25.71881112	2.660382523	Up	5.35E-14	1.80E-13	AAEL005259-PA OS=Aedes aegypti GN=AAEL005259 PE=4 SV=1

CCG004648.1	1967	73	373	1.594708	6.629098747	2.055520465	Up	0	0	AAEL001996-PA OS=Aedes aegypti GN=AAEL001996 PE=4 SV=1
CCG023137.1	2214	80	373	1.55265527	5.889538047	1.923416929	Up	0	0	AAEL002514-PA OS=Aedes aegypti GN=AAEL002514 PE=4 SV=1
CCG006032.1	1473	90	373	2.62544203	8.852299549	1.753491928	Up	0	0	AAEL002864-PA OS=Aedes aegypti GN=AAEL002864 PE=4 SV=1
CCG017755.1	1066	101	373	4.07124127	12.23211748	1.587133541	Up	0	0	AAEL008691-PA OS=Aedes aegypti GN=AAEL008691 PE=4 SV=1
CCG008557.1	710	102	373	6.1731168	18.36540456	1.572919682	Up	0	0	Ubiquitin conjugation factor E4A OS=Culex quinquefasciatus GN=CpipJ_CPIJ001165 PE=4 SV=1
CCG014006.1	1332	726	373	23.4204409	9.789367294	-1.258480714	Down	2.27E-45	1.04E-44	AAEL006630-PA OS=Aedes aegypti GN=AAEL006630 PE=4 SV=1
CCG015692.1	2358	1146	373	20.8835097	5.529871601	-1.917046305	Down	7.82E-130	4.34E-129	AAEL014316-PA (Fragment) OS=Aedes aegypti GN=AAEL014316 PE=4 SV=1
CCG027807.1	465	40	374	3.69632126	28.1169796	2.927279569	Up	6.86E-14	2.25E-13	AAEL000408-PA OS=Aedes aegypti GN=AAEL000408 PE=4 SV=1
CCG021152.1	849	76	374	3.84652512	15.39975914	2.00128015	Up	0	0	AAEL011171-PA OS=Aedes aegypti GN=AAEL011171 PE=3 SV=1
CCG010949.1	1644	134	374	3.50239929	7.952795324	1.183118473	Up	0	0	AAEL002560-PA OS=Aedes aegypti GN=AAEL002560 PE=4 SV=1
CCG010098.1	996	52	375	2.2433998	13.1620018	2.552620271	Up	0	0	Ribosomal protein S6 kinase, 90kD, polypeptide OS=Culex quinquefasciatus GN=CpipJ_CPIJ012176 PE=4 SV=1
CCG028220.1	327	102	375	13.4034035	40.08976694	1.580634647	Up	0	0	-
CCG002793.1	443	106	375	10.2816972	29.59222074	1.525139535	Up	0	0	Shmp sm protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001194 PE=4 SV=1
CCG021868.1	1107	110	375	4.26980199	11.84223468	1.471700276	Up	0	0	AAEL001326-PA OS=Aedes aegypti GN=AAEL001326 PE=4 SV=1
CCG006255.1	3192	138	375	1.85771409	4.10694041	1.144535532	Up	1.49E-13	4.45E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013304 PE=4 SV=1

CCG006264.1	468	144	375	13.2214568	28.01143972	1.083134988	Up	7.62E-14	2.46E-13	Peptidyl-prolyl cis-trans isomerase OS=Aedes aegypti GN=AAEL004779 PE=3 SV=1
CCG010078.1	483	649	375	57.7378008	27.14151923	-1.089014679	Down	1.21E-32	5.15E-32	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ019370 PE=4 SV=1
CCG016315.1	1173	1451	375	53.1535251	11.17591968	-2.249771815	Down	1.18E-201	7.06E-201	AAEL0008963-PA OS=Aedes aegypti GN=AAEL008963 PE=3 SV=1
CCG025054.1	2522	6859	375	116.863366	5.197999124	-4.490722551	Down	0	0	AAEL0005569-PA OS=Aedes aegypti GN=AAEL005569 PE=4 SV=1
CCG023622.1	849	10	376	0.50612173	15.4821108	4.934973961	Up	5.43E-07	1.01E-06	-
CCG004904.1	1080	26	376	1.03445657	12.17065932	3.556462337	Up	4.60E-12	1.05E-11	AAEL000411-PA OS=Aedes aegypti GN=AAEL000411 PE=4 SV=1
CCG001507.1	1869	39	376	0.89663973	7.032804744	2.971499837	Up	7.33E-14	2.38E-13	AAEL0003178-PA OS=Aedes aegypti GN=AAEL003178 PE=4 SV=1
CCG026487.1	1929	41	376	0.91330177	6.814054985	2.899350051	Up	1.02E-13	3.19E-13	Zinc finger protein 394 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005856 PE=4 SV=1
CCG011159.1	3309	106	376	1.37648591	3.972291347	1.528981601	Up	0	0	AAEL0006614-PA OS=Aedes aegypti GN=AAEL006614 PE=4 SV=1
CCG020707.1	3213	125	376	1.67171392	4.090977923	1.291117771	Up	2.35E-14	8.22E-14	AAEL0005429-PA OS=Aedes aegypti GN=AAEL005429 PE=4 SV=1
CCG012894.1	1533	139	376	3.89614684	8.5742414	1.137960983	Up	1.46E-13	4.37E-13	AAEL0009499-PA OS=Aedes aegypti GN=AAEL009499 PE=4 SV=1
CCG016477.1	489	1	377	0.08787267	26.95147309	8.260733917	Up	0.000587	0.00091598	AAEL013262-PA OS=Aedes aegypti GN=AAEL013262 PE=4 SV=1
CCG000857.1	2448	38	377	0.66701385	5.383688866	3.012806404	Up	4.26E-14	1.45E-13	AAEL012906-PA OS=Aedes aegypti GN=AAEL012906 PE=4 SV=1
CCG024187.1	1960	40	377	0.87693336	6.724117522	2.938805822	Up	6.86E-14	2.24E-13	AAEL0008769-PA OS=Aedes aegypti GN=AAEL008769 PE=3 SV=1
CCG026931.1	1278	81	377	2.72343388	10.31241811	1.920883914	Up	4.46E-14	1.51E-13	AAEL010728-PA OS=Aedes aegypti GN=AAEL010728 PE=4 SV=1
CCG027145.1	1780	100	377	2.41403003	7.404084462	1.616877727	Up	0	0	AAEL014348-PB OS=Aedes aegypti GN=CASPS8 PE=4 SV=1

CCG019120.2	1152	122	377	4.55061426	11.44033884	1.32999658	Up	8.04E-14	2.58E-13	AAEL000461-PA OS=Aedes aegypti GN=AAEL000461 PE=4 SV=1
CCG025975.1	1590	1395	377	37.6998615	8.288849272	-2.18531549	Down	5.23E-187	3.09E-186	AAEL013611-PA OS=Aedes aegypti GN=AAEL013611 PE=4 SV=1
CCG008307.1	2313	5336	377	99.1294872	5.697911951	-4.120809034	Down	0	0	Myosin light chain kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005740 PE=4 SV=1
CCG019130.1	734	1	378	0.05854187	18.00303627	8.264555628	Up	0.000587	0.00091564	AAEL009973-PA OS=Aedes aegypti GN=AAEL009973 PE=4 SV=1
CCG001120.1	791	29	378	1.57537586	16.70572518	3.406574633	Up	6.66E-13	1.69E-12	AAEL006819-PA OS=Aedes aegypti GN=AAEL006819 PE=4 SV=1
CCG006386.1	1379	45	378	1.40220309	9.582471806	2.772702532	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008646 PE=4 SV=1
CCG027535.1	1551	54	378	1.49604492	8.519812134	2.509668126	Up	0	0	AAEL001785-PA OS=Aedes aegypti GN=AAEL001785 PE=4 SV=1
CCG006131.1	1134	58	378	2.19774657	11.65275892	2.406574633	Up	0	0	AAEL008544-PA OS=Aedes aegypti GN=AAEL008544 PE=4 SV=1
CCG014579.1	1736	84	378	2.07918071	7.611882846	1.872238205	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011278 PE=4 SV=1
CCG028197.1	900	85	378	4.05825271	14.68247624	1.855164692	Up	0	0	Protein kinase C OS=Aedes aegypti GN=AAEL001108 PE=3 SV=1
CCG010492.1	1107	86	378	3.33820883	11.93697256	1.838290873	Up	0	0	AAEL005637-PA OS=Aedes aegypti GN=AAEL005637 PE=4 SV=1
CCG021584.1	1010	96	378	4.084252	13.08339467	1.679593127	Up	0	0	AAEL002836-PA (Fragment) OS=Aedes aegypti GN=AAEL002836 PE=4 SV=1
CCG023666.1	984	98	378	4.27950609	13.42909413	1.649845784	Up	0	0	AAEL009589-PA OS=Aedes aegypti GN=AAEL009589 PE=4 SV=1
CCG000055.1	737	150	378	8.74553622	17.92975389	1.035736938	Up	0	0	AAEL011710-PA OS=Aedes aegypti GN=AAEL011710 PE=4 SV=1
CCG003501.1	744	1307	378	75.4858107	17.76105997	-2.087487798	Down	2.26E-165	1.31E-164	AAEL014944-PA OS=Aedes aegypti GN=SGP2D7 PE=2 SV=1

CCG005099.1	405	1571	378	166.680131	32.62772499	-2.352911837	Down	1.59E-230	9.65E-230	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004235 PE=4 SV=1
CCG023160.1	1119	45	379	1.72800541	11.84020277	2.776514146	Up	0	0	AAEL013503-PA OS=Aedes aegypti GN=AAEL013503 PE=4 SV=1
CCG013116.1	606	49	379	3.47445049	21.86334471	2.653657398	Up	0	0	AAEL005704-PA OS=Aedes aegypti GN=AAEL005704 PE=4 SV=1
CCG001584.1	4632	66	379	0.61226306	2.860359865	2.223973123	Up	0	0	AAEL002922-PA OS=Aedes aegypti GN=AAEL002922 PE=4 SV=1
CCG014808.1	1879	75	379	1.71513044	7.051190472	2.039548551	Up	0	0	NADPH:adrenodoxin oxidoreductase, mitochondrial OS=Aedes aegypti GN=AAEL006733 PE=3 SV=1
CCG005376.1	1089	77	379	3.03826406	12.16637915	2.001580701	Up	0	0	AAEL011989-PA OS=Aedes aegypti GN=AAEL011989 PE=4 SV=1
CCG023379.1	1215	79	379	2.7939169	10.90468057	1.964586494	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_02694 PE=4 SV=1
CCG007383.2	1119	80	379	3.07200962	11.84020277	1.946439147	Up	0	0	AGAP001896-PA OS=Anopheles gambiae GN=AGAP001896 PE=4 SV=1
CCG008694.1	1788	146	379	3.50871435	7.410059786	1.078542683	Up	9.99E-15	3.57E-14	AAEL009899-PA OS=Aedes aegypti GN=AAEL009899 PE=4 SV=1
CCG005111.1	438	2435	379	238.884255	30.24928515	-2.981338815	Down	0	0	AAEL001164-PA OS=Aedes aegypti GN=AAEL001164 PE=4 SV=1
CCG019237.1	984	54	380	2.35809519	13.50014753	2.51728131	Up	0	0	AAEL005813-PA OS=Aedes aegypti GN=AAEL005813 PE=4 SV=1
CCG020342.1	1797	66	380	1.57818725	7.392401321	2.227774693	Up	0	0	AAEL012156-PA OS=Aedes aegypti GN=AAEL012156 PE=4 SV=1
CCG023531.2	2283	687	380	12.9304458	5.818723247	-1.151997477	Down	1.37E-37	6.02E-37	AAEL007180-PA OS=Aedes aegypti GN=AAEL007180 PE=4 SV=2
CCG005933.1	735	6	381	0.35077334	18.12122918	5.690997891	Up	8.38E-06	1.47E-05	AAEL005000-PA OS=Aedes aegypti GN=AAEL005000 PE=4 SV=1
CCG022938.1	1866	22	381	0.50660995	7.137783199	3.816528773	Up	8.38E-11	1.80E-10	AAEL014291-PA OS=Aedes aegypti GN=AAEL014291 PE=4 SV=1

CCG000130.1	330	40	381	5.20845268	40.36091955	2.954032296	Up	6.86E-14	2.24E-13	AAEL014055-PA OS=Aedes aegypti GN=AAEL014055 PE=4 SV=1
CCG023654.1	966	45	381	2.00169571	13.78789177	2.784107295	Up	0	0	Cuiclicron OS=Culex quinquefasciatus GN=CpipJ_CPIJ005389 PE=4 SV=1
CCG013164.1	1717	49	381	1.22627664	7.757194788	2.661250547	Up	0	0	AAEL008392-PA OS=Aedes aegypti GN=AAEL008392 PE=4 SV=1
CCG019080.1	1257	81	381	2.76893278	10.59594547	1.936110388	Up	4.46E-14	1.51E-13	AAEL003744-PA (Fragment) OS=Aedes aegypti GN=AAEL003744 PE=4 SV=1
CCG024032.1	1230	105	381	3.66814808	10.82853939	1.561714874	Up	0	0	AAEL004682-PA OS=Aedes aegypti GN=AAEL004682 PE=4 SV=1
CCG025554.1	1002	112	381	4.80300427	13.29251841	1.468605469	Up	0	0	AAEL007996-PA OS=Aedes aegypti GN=AAEL007996 PE=4 SV=1
CCG014946.1	978	1584	381	69.595153	13.61871518	-2.353396229	Down	1.78E-232	1.09E-231	tRNA pseudouridine synthase OS=Aedes aegypti GN=AAEL003683 PE=3 SV=1
CCG009321.1	1332	19	382	0.61293165	10.02557187	4.031814518	Up	7.73E-10	1.60E-09	AAEL010588-PA (Fragment) OS=Aedes aegypti GN=AAEL010588 PE=4 SV=1
CCG007463.1	1868	52	382	1.19615964	7.148855314	2.579302314	Up	0	0	Gastrula zinc finger protein XLCGF62.1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007367 PE=4 SV=1
CCG009227.1	5034	63	382	0.53776188	2.652773486	2.302462108	Up	0	0	AAEL003012-PA OS=Aedes aegypti GN=AAEL003012 PE=4 SV=1
CCG025348.1	1027	7610	382	318.402805	13.00298123	-4.613938707	Down	0	0	-
CCG002383.1	1386	3	383	0.09300808	9.660187593	6.698551285	Up	5.28E-05	8.84E-05	AAEL010782-PA OS=Aedes aegypti GN=AAEL010782 PE=3 SV=1
CCG019134.1	1974	17	383	0.37005344	6.782684906	4.196050944	Up	3.39E-09	6.89E-09	AAEL002281-PA OS=Aedes aegypti GN=AAEL002281 PE=4 SV=1
CCG007063.1	1905	24	383	0.54135099	7.028356957	3.698551285	Up	1.93E-11	4.23E-11	AAEL002250-PA OS=Aedes aegypti GN=AAEL002250 PE=4 SV=1
CCG012794.1	1359	25	383	0.79046605	9.85211185	3.639657596	Up	9.32E-12	2.08E-11	AAEL010331-PA OS=Aedes aegypti GN=AAEL010331 PE=4 SV=1

CCG027870.1	564	46	383	3.50462374	23.73939717	2.75995183	Up	3.40E-14	1.18E-13	-
CCG005955.1	840	114	383	5.83160684	15.93930953	1.450623772	Up	8.48E-14	2.70E-13	NADPH17AD oxidoreductase OS=Culex quinquefasciatus GN=CpipJ_CPIJ003677 PE=4 SV=1
CCG018494.1	7563	1788	383	10.1586521	1.770331879	-2.520617235	Down	1.84E-284	1.15E-283	Tyrosine-protein kinase receptor (Fragment) OS=Aedes aegypti GN=AAEL007742 PE=3 SV=1
CCG011840.1	2091	11	384	0.22604834	6.4198844	4.827844086	Up	2.67E-07	5.03E-07	AAEL004869-PA (Fragment) OS=Aedes aegypti GN=AAEL004869 PE=4 SV=1
CCG022196.1	967	40	384	1.77744507	13.88208716	2.96534761	Up	6.86E-14	2.24E-13	AAEL008320-PA OS=Aedes aegypti GN=AAEL008320 PE=4 SV=1
CCG010014.1	2499	44	384	0.75656996	5.371740008	2.827844086	Up	3.91E-14	1.34E-13	AAEL009815-PA OS=Aedes aegypti GN=AAEL009815 PE=4 SV=1
CCG001165.1	2595	61	384	1.01007854	5.173016678	2.356538367	Up	1.75E-14	6.17E-14	AAEL002195-PA (Fragment) OS=Aedes aegypti GN=AAEL002195 PE=4 SV=1
CCG026438.1	1300	65	384	2.14848673	10.32613714	2.264907892	Up	0	0	AAEL003526-PA OS=Aedes aegypti GN=AAEL003526 PE=4 SV=1
CCG023932.1	860	67	384	3.34764212	15.60927707	2.221186514	Up	0	0	AAEL010053-PA OS=Aedes aegypti GN=AAEL010053 PE=4 SV=1
CCG003203.1	1113	71	384	2.74110616	12.06107662	2.137528585	Up	0	0	AAEL005064-PA OS=Aedes aegypti GN=CLIPB5 PE=3 SV=1
CCG009136.1	1532	83	384	2.32799476	8.762387912	1.912236273	Up	1.32E-13	4.02E-13	AAEL005635-PA OS=Aedes aegypti GN=AAEL005635 PE=4 SV=1
CCG011630.1	2385	94	384	1.69356606	5.628502424	1.732686853	Up	0	0	AAEL009919-PA OS=Aedes aegypti GN=AAEL009919 PE=4 SV=1
CCG027707.1	1340	123	384	3.94423683	10.01789424	1.344761199	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_17122 PE=4 SV=1
CCG004357.1	714	130	384	7.82362115	18.80109003	1.264907892	Up	0	0	FAM105 OS=Culex quinquefasciatus GN=CpipJ_CPIJ016405 PE=4 SV=1

CCG015468.1	3642	627	384	7.39759023	3.685880912	-1.005045928	Down	6.93E-28	2.88E-27	Serine-enriched protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005696 PE=4 SV=1
CCG016960.1	3861	1582	384	17.6063507	3.476813851	-2.34025818	Down	1.31E-230	7.97E-230	AAEL000037-PA OS=Aedes aegypti GN=CLIPB35 PE=3 SV=1
CCG025026.1	2448	1607	384	28.2076648	5.483651258	-2.362878509	Down	5.65E-237	3.45E-236	microtubule-associated protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013260 PE=4 SV=1
CCG014070.1	807	68	385	3.62074592	16.67774047	2.203564998	Up	0	0	Fat body protein 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ018909 PE=3 SV=1
CCG012221.1	1820	16	386	0.37775591	7.414227931	4.294770241	Up	7.09E-09	1.42E-08	AAEL002058-PA OS=Aedes aegypti GN=AAEL002838 PE=4 SV=1
CCG026687.1	3939	30	386	0.32726378	3.425715876	3.387879645	Up	3.71E-13	9.95E-13	AAEL000658-PA OS=Aedes aegypti GN=AAEL006658 PE=4 SV=1
CCG004327.1	651	54	386	3.56430978	20.72794905	2.539882739	Up	0	0	AAEL013578-PA OS=Aedes aegypti GN=AAEL013578 PE=4 SV=1
CCG011014.1	780	156	386	8.59394692	17.29986517	1.009368022	Up	1.59E-13	4.71E-13	AAEL013225-PA OS=Aedes aegypti GN=AAEL013225 PE=4 SV=1
CCG026050.1	390	770	386	84.8376812	34.59973034	-1.293944394	Down	4.43E-50	2.07E-49	AAEL009810-PA OS=Aedes aegypti GN=AAEL009810 PE=2 SV=1
CCG006328.1	1836	31	387	0.72552384	7.368656378	3.34430665	Up	2.40E-13	6.78E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009123 PE=4 SV=1
CCG018315.1	785	55	387	3.01061835	17.23420778	2.517143246	Up	0	0	Mediator complex OS=Culex quinquefasciatus GN=CpipJ_CPIJ006001 PE=4 SV=1
CCG010986.1	1700	127	387	3.21009194	7.958148889	1.309818273	Up	1.57E-13	4.67E-13	S-methyl-5-riboadenosine phosphorylase OS=Culex quinquefasciatus GN=CpipJ_CPIJ002278 PE=3 SV=1
CCG023353.1	20046	863	387	1.8498893	0.674890408	-1.454713789	Down	1.30E-66	6.37E-66	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016863 PE=4 SV=1

CCG016626.1	6855	1379	387	8.64409395	1.973574487	-2.130903782	Down	4.72E-179	2.76E-178	AAEL007465-PA OS=Aedes aegypti GN=AAEL007465 PE=4 SV=1
CCG006583.1	676	2463	387	156.559847	20.01309632	-2.967697953	Down	0	0	AAEL007283-PA OS=Aedes aegypti GN=AAEL007283 PE=4 SV=1
CCG002975.1	1488	15	388	0.43316265	9.115464642	4.39533545	Up	1.48E-08	2.93E-08	AAEL009709-PA OS=Aedes aegypti GN=AAEL009709 PE=4 SV=1
CCG013525.1	2674	17	388	0.27318081	5.072479951	4.214763205	Up	3.39E-09	6.89E-09	Polyprotein OS=Antheraea mylitta PE=4 SV=1
CCG005988.1	1798	32	388	0.76475612	7.543832807	3.302226046	Up	1.77E-13	5.17E-13	AAEL004125-PA OS=Aedes aegypti GN=AAEL004125 PE=4 SV=1
CCG020412.1	1941	94	388	2.08096602	6.988053265	1.747637194	Up	0	0	AAEL002594-PA (Fragment) OS=Aedes aegypti GN=AAEL002594 PE=4 SV=1
CCG027007.1	1049	114	388	4.66973284	12.93023011	1.469336032	Up	8.48E-14	2.70E-13	AAEL007647-PA OS=Aedes aegypti GN=AAEL007647 PE=4 SV=1
CCG003837.1	1458	670	388	19.7460372	9.303025643	-1.085791239	Down	1.71E-33	7.35E-33	AAEL001699-PA OS=Aedes aegypti GN=AAEL001699 PE=4 SV=1
CCG006390.1	777	713	388	39.4304	17.45664271	-1.17553222	Down	3.52E-40	1.57E-39	Protein OS=Culex quinquefasciatus
CCG019541.1	1014	38	389	1.61030564	13.41101545	3.058012035	Up	4.26E-14	1.45E-13	AAEL008716-PA OS=Aedes aegypti GN=AAEL008716 PE=4 SV=1
CCG023905.2	663	52	389	3.37017526	20.5109648	2.605499831	Up	0	0	AAEL001871-PA OS=Aedes aegypti GN=AAEL001871 PE=4 SV=1
CCG008308.1	663	59	389	3.8238527	20.5109648	2.423296499	Up	0	0	Dacapo OS=Aedes albopictus PE=2 SV=1
CCG006921.1	830	67	389	3.46864123	16.38405984	2.239850358	Up	0	0	AAEL004448-PA OS=Aedes aegypti GN=AAEL004448 PE=4 SV=1
CCG019035.1	927	73	389	3.38380866	14.66965444	2.11611499	Up	0	0	AAEL004900-PA OS=Aedes aegypti GN=AAEL004900 PE=3 SV=1
CCG024275.1	1482	74	389	2.14558729	9.175957938	2.096486183	Up	0	0	AAEL014397-PA OS=Aedes aegypti GN=AAEL014397 PE=4 SV=1
CCG004930.1	1203	80	389	2.85750521	11.30404793	1.984011454	Up	0	0	AAEL012040-PA OS=Aedes aegypti GN=AAEL012040 PE=4 SV=1
CCG020971.1	378	89	389	10.1172126	35.97558112	1.830206118	Up	0	0	AAEL009372-PA OS=Aedes aegypti GN=AAEL009372 PE=4 SV=1

CCG003915.1	593	93	389	6.73892971	22.93215795	1.766780738	Up	4.04E-14	1.38E-13	AAEL005705-PA OS=Aedes aegypti GN=AAEL005705 PE=4 SV=1
CCG020634.1	936	102	389	4.68259928	14.52860007	1.633514207	Up	0	0	AAEL001405-PA OS=Aedes aegypti GN=AAEL001405 PE=4 SV=1
CCG026690.1	1106	744	389	28.9054996	12.29545178	-1.233219262	Down	4.85E-45	2.21E-44	AAEL010592-PA OS=Aedes aegypti GN=AAEL010592 PE=4 SV=1
CCG013552.1	1182	22	390	0.79977509	11.5344568	3.850211899	Up	8.38E-11	1.80E-10	AAEL000287-PA (Fragment) OS=Aedes aegypti GN=AAEL000287 PE=4 SV=1
CCG004315.1	1083	32	390	1.26965051	12.58885313	3.309643518	Up	1.77E-13	5.17E-13	AAEL006547-PA OS=Aedes aegypti GN=AAEL006547 PE=2 SV=1
CCG025639.1	2271	46	390	0.87036891	6.00340288	2.786081562	Up	3.40E-14	1.18E-13	AAEL014110-PA OS=Aedes aegypti GN=AAEL014110 PE=4 SV=1
CCG019372.1	597	53	390	3.81473356	22.83706523	2.581723063	Up	0	0	AAEL007877-PA OS=Aedes aegypti GN=AAEL007877 PE=4 SV=1
CCG014283.1	867	108	390	5.3526313	15.7251764	1.554756015	Up	0	0	AAEL015082-PA (Fragment) OS=Aedes aegypti GN=AAEL015082 PE=4 SV=1
CCG027490.1	2681	111	390	1.77905279	5.085314413	1.515227651	Up	0	0	AAEL002248-PA (Fragment) OS=Aedes aegypti GN=AAEL002248 PE=4 SV=1
CCG002492.1	922	114	390	5.31296068	14.78712358	1.476753503	Up	8.48E-14	2.70E-13	-
CCG010836.1	1275	132	390	4.44863135	10.69311995	1.265249398	Up	1.46E-13	4.36E-13	AAEL003125-PA OS=Aedes aegypti GN=AAEL003125 PE=3 SV=1
CCG015275.1	1221	137	390	4.82133795	11.16603435	1.211611435	Up	2.90E-13	8.03E-13	AAEL005103-PA OS=Aedes aegypti GN=AAEL005103 PE=4 SV=1
CCG007206.1	1956	31	391	0.68101318	6.988080889	3.359141691	Up	2.40E-13	6.76E-13	Putative uncharacterized protein OS=Daphnia pulex GN=DAPPUDRAFT_311987 PE=4 SV=1
CCG008311.1	2844	64	391	0.96697012	4.806148459	2.313338001	Up	0	0	AAEL003562-PA OS=Aedes aegypti GN=AAEL003562 PE=4 SV=1
CCG017062.1	4106	76	391	0.79534823	3.328954266	2.065410488	Up	0	0	AAEL009531-PA OS=Aedes aegypti GN=AAEL009531 PE=4 SV=1

CCG020541.1	2988	102	391	1.46683833	4.574526847	1.640912659	Up	0	0	AAEL013167-PA OS=Aedes aegypti GN=AAEL013167 PE=4 SV=1
CCG021399.1	1187	132	391	4.77843721	11.51532116	1.268943882	Up	1.46E-13	4.36E-13	-
CCG014038.1	774	145	391	8.04988568	17.65980132	1.133428911	Up	0	0	AAEL013197-PA OS=Aedes aegypti GN=AAEL013197 PE=4 SV=1
CCG008236.1	1361	905	391	28.5728213	10.04311993	-1.508435981	Down	1.42E-73	7.09E-73	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010714 PE=4 SV=1
CCG028232.1	1140	2492	391	93.9303322	11.99007563	-2.969750352	Down	0	0	AAEL012556-PA OS=Aedes aegypti GN=AAEL012556 PE=4 SV=1
CCG022827.1	753	51	392	2.91030075	18.19873107	2.644597706	Up	0	0	Phosphatase and actin regulator OS=Culex quinquefasciatus GN=CpipJ_CPIJ018341 PE=4 SV=1
CCG003477.1	1488	58	392	1.67489557	9.209438505	2.459042053	Up	0	0	AAEL013014-PA OS=Aedes aegypti GN=AAEL013014 PE=4 SV=1
CCG019518.1	405	139	392	14.7476373	33.83615925	1.198081975	Up	1.46E-13	4.37E-13	AAEL008044-PA OS=Aedes aegypti GN=AAEL008044 PE=4 SV=1
CCG019446.1	3393	150	392	1.8996346	4.038798849	1.088204357	Up	0	0	Pol-like protein OS=Aedes aegypti PE=4 SV=1
CCG003793.1	1596	792	392	21.3233269	8.586243418	-1.312333572	Down	1.44E-52	6.77E-52	AAEL001920-PA OS=Aedes aegypti GN=AAEL001920 PE=4 SV=1
CCG027245.1	1208	5	393	0.17785486	11.37301554	5.998770611	Up	1.59E-05	2.76E-05	AAEL007829-PA (Fragment) OS=Aedes aegypti GN=AAEL007829 PE=4 SV=1
CCG003177.2	1221	32	393	1.12615193	11.25192692	3.320698706	Up	1.77E-13	5.16E-13	AAEL005406-PA OS=Aedes aegypti GN=AAEL005406 PE=4 SV=1
CCG018648.1	1125	49	393	1.87157066	12.21209135	2.705988862	Up	0	0	AAEL013921-PA OS=Aedes aegypti GN=AAEL013921 PE=4 SV=1
CCG017323.2	777	57	393	3.15221991	17.68159945	2.487808692	Up	0	0	-
CCG019389.1	1512	123	393	3.49555381	9.086377494	1.378184201	Up	0	0	tRNA pseudouridine synthase OS=Aedes aegypti GN=AAEL007354 PE=3 SV=1
CCG019331.1	3705	129	393	1.49611222	3.708124905	1.309471451	Up	0	0	AAEL007127-PA (Fragment) OS=Aedes aegypti GN=AAEL007127 PE=4 SV=1

CCG017527.1	15291	997	393	2.80170201	0.898476409	-1.640750988	Down	2.71E-91	1.41E-90	AAEL010027-PA OS=Aedes aegypti GN=AAEL010027 PE=4 SV=1
CCG021649.1	1388	2432	393	75.2899096	9.898128798	-2.927228807	Down	0	0	AAEL000354-PA OS=Aedes aegypti GN=AAEL000354 PE=4 SV=1
CCG005228.1	432	18	394	1.79040561	31.88324317	4.154440022	Up	1.62E-09	3.33E-09	AAEL000555-PA OS=Aedes aegypti GN=AAEL008535 PE=4 SV=1
CCG020843.1	1347	62	394	1.97782	10.22536084	2.370168713	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_17290 PE=4 SV=1
CCG019736.1	708	87	394	5.28017925	19.45418227	1.881421527	Up	0	0	-
CCG016861.1	1450	90	394	2.66708698	9.49900762	1.832511927	Up	0	0	AAEL003649-PA OS=Aedes aegypti GN=AAEL003649 PE=4 SV=1
CCG019574.1	1521	92	394	2.5990898	9.055595692	1.800803067	Up	1.44E-14	5.12E-14	AAEL001787-PA OS=Aedes aegypti GN=AAEL001787 PE=4 SV=1
CCG013767.1	1044	101	394	4.15703371	13.19306614	1.666153541	Up	0	0	AAEL011338-PA OS=Aedes aegypti GN=AAEL011338 PE=4 SV=1
CCG021562.1	876	114	394	5.59195176	15.72324321	1.491475009	Up	8.48E-14	2.70E-13	Cleavage and polyadenylation specificity factor subunit 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009192 PE=4 SV=1
CCG015596.1	780	784	394	43.1900922	17.6584116	-1.290344821	Down	9.59E-51	4.48E-50	AAEL003716-PA OS=Aedes aegypti GN=AAEL003716 PE=2 SV=1
CCG023125.1	696	25	395	1.54345311	19.83982662	3.684165857	Up	9.32E-12	2.08E-11	AAEL011866-PA OS=Aedes aegypti GN=AAEL011866 PE=4 SV=1
CCG002578.1	717	62	395	3.71565348	19.25874383	2.373825736	Up	0	0	AAEL014041-PA OS=Aedes aegypti GN=AAEL014041 PE=4 SV=1
CCG002295.1	3477	77	395	0.95158745	3.971388934	2.061235506	Up	0	0	AAEL003438-PA OS=Aedes aegypti GN=AAEL003438 PE=4 SV=1
CCG005517.1	735	87	395	5.08621348	18.78710112	1.885078551	Up	0	0	AAEL006172-PA OS=Aedes aegypti GN=AAEL006172 PE=4 SV=1
CCG003520.1	1406	143	395	4.37032151	9.8211375	1.16815071	Up	0	0	AAEL014512-PA (Fragment) OS=Aedes aegypti GN=AAEL014512 PE=4 SV=1
CCG017777.1	615	158	395	11.039379	22.45287695	1.024241299	Up	0	0	Male specific 1 protein OS=Aedes aegypti GN=DSXM1 PE=2 SV=1

CCG016414.1	3621	657	395	7.796497	3.81345466	-1.031727513	Down	2.32E-30	9.78E-30	AAEL004678-PA OS=Aedes aegypti GN=AAEL004678 PE=4 SV=1
CCG010647.1	389	2	396	0.22092409	35.58734602	7.331669824	Up	9.36E-05	0.00015353	Chromosome undetermined SCAF14565, whole genome shotgun sequence OS=Tetraodon nigroviridis GN=GSTENG00016696001 PE=3 SV=1
CCG000834.1	638	116	396	7.81267902	21.69824076	1.473688829	Up	1.16E-13	3.59E-13	AAEL009089-PA OS=Aedes aegypti GN=AAEL009089 PE=4 SV=1
CCG023129.1	2448	37	397	0.64946086	5.669295702	3.125855035	Up	5.17E-14	1.74E-13	Ribonuclease P/MKP protein subunit POP1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007891 PE=4 SV=1
CCG017197.1	1395	91	397	2.80304362	9.948699555	1.827513761	Up	0	0	AAEL002329-PA OS=Aedes aegypti GN=AAEL002329 PE=4 SV=1
CCG026724.1	1074	126	397	5.04114205	12.92219356	1.358028477	Up	1.34E-13	4.06E-13	AAEL005093-PA OS=Aedes aegypti GN=CLIPB46 PE=3 SV=1
CCG000376.1	792	136	397	7.3786413	17.52327762	1.24784556	Up	1.70E-13	4.99E-13	AAEL005036-PA OS=Aedes aegypti GN=AAEL005036 PE=4 SV=1
CCG013950.1	23081	689	397	1.28270643	0.60129266	-1.093051772	Down	9.41E-35	4.07E-34	AAEL008894-PA OS=Aedes aegypti GN=AAEL008894 PE=4 SV=1
CCG025148.1	23271	880	397	1.62491369	0.596383304	-1.446051313	Down	2.71E-67	1.33E-66	AAEL002172-PA OS=Aedes aegypti GN=AAEL002172 PE=4 SV=1
CCG013374.1	969	1702	397	75.4741881	14.32243125	-2.397706921	Down	2.49E-255	1.54E-254	AAEL005337-PA OS=Aedes aegypti GN=AAEL005337 PE=4 SV=1
CCG005495.1	1488	35	398	1.01071284	9.350399298	3.209654807	Up	6.39E-14	2.11E-13	Stm1 cell tumor OS=Culex quinquefasciatus GN=CpipJ_CPIJ003969 PE=4 SV=1
CCG017044.1	825	53	398	2.76047992	16.86472019	2.61101737	Up	0	0	AAEL008790-PA OS=Aedes aegypti GN=AAEL008790 PE=4 SV=1
CCG011228.1	864	73	398	3.63054471	16.10346546	2.149113265	Up	0	0	AAEL010343-PA (Fragment) OS=Aedes aegypti GN=AAEL010343 PE=4 SV=1

CCG025127.1	429	3798	398	380.417371	32.43215421	-3.552086366	Down	0	0	Putative salivary odorant binding protein 1 OS=Culex quinquefasciatus PE=2 SV=1
CCG007264.1	1116	70	399	2.69523425	12.49852368	2.213275123	Up	0	0	AAEL012414-PA OS=Aedes aegypti GN=AAEL012414 PE=4 SV=1
CCG023888.1	7149	1218	399	7.32090317	1.951091402	-1.907740278	Down	5.71E-137	3.20E-136	AAEL001803-PA (Fragment) OS=Aedes aegypti GN=AAEL001803 PE=4 SV=1
CCG017693.1	1008	1928	399	82.1881432	13.83765122	-2.570331196	Down	0	0	AAEL010783-PA OS=Aedes aegypti GN=AAEL010783 PE=3 SV=1
CCG026736.1	2743	84	400	1.31587959	5.097816518	1.953851971	Up	0	0	AAEL006317-PA OS=Aedes aegypti GN=AAEL006317 PE=4 SV=1
CCG008321.1	1437	99	400	2.96033662	9.730905156	1.716812774	Up	0	0	AAEL013521-PA OS=Aedes aegypti GN=AAEL013521 PE=3 SV=1
CCG003656.1	2193	128	400	2.50803741	6.376338673	1.346169394	Up	4.15E-14	1.42E-13	AAEL011047-PA (Fragment) OS=Aedes aegypti GN=AAEL011047 PE=4 SV=1
CCG017437.1	672	986	400	63.0478547	20.80849808	-1.599274443	Down	4.60E-87	2.37E-86	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001349 PE=4 SV=1
CCG022081.1	1104	1301	400	50.6373412	12.66604231	-1.999235853	Down	1.36E-155	7.78E-155	AAEL001261-PA OS=Aedes aegypti GN=AAEL001261 PE=4 SV=1
CCG014058.1	2428	2250	400	39.8195646	5.759188925	-2.789539893	Down	0	0	AAEL008028-PA OS=Aedes aegypti GN=AAEL008028 PE=4 SV=1
CCG002538.1	1233	29	401	1.01064258	11.36923681	3.491790635	Up	6.66E-13	1.69E-12	AAEL001939-PA OS=Aedes aegypti GN=AAEL001939 PE=4 SV=1
CCG023116.1	7988	83	401	0.44648072	1.754915997	1.974732199	Up	1.32E-13	4.02E-13	AAEL014263-PA OS=Aedes aegypti GN=AAEL014263 PE=4 SV=1
CCG026306.1	830	116	401	6.00540869	16.88948071	1.491790635	Up	1.16E-13	3.59E-13	Phosphataseppt regulatory subunit OS=Culex quinquefasciatus GN=CpipJ_CPIJ003122 PE=4 SV=1
CCG014446.1	1152	134	401	4.99821566	12.16863627	1.28368244	Up	0	0	AAEL014712-PA OS=Aedes aegypti GN=AAEL014712 PE=4 SV=1
CCG012745.1	464	999	401	92.5145795	30.21178661	-1.614569238	Down	2.13E-89	1.11E-88	AAEL004972-PA OS=Aedes aegypti GN=AAEL004972 PE=4 SV=1

CCG024624.1	398	1	402	0.10796416	35.30961624	8.353364895	Up	0.000587	0.00091541	Chromosome undetermined SCAF14565, whole genome shotgun sequence OS=Tetraodon nigroviridis GN=GSTENG00016696001 PE=3 SV=1
CCG026310.1	903	81	402	3.85442802	15.56282089	2.013514892	Up	4.46E-14	1.51E-13	AAEL003151-FA OS=Aedes aegypti GN=AAEL003151 PE=4 SV=1
CCG021465.1	1149	86	402	3.21618553	12.23083313	1.92710014	Up	0	0	AAEL012211-FA OS=Aedes aegypti GN=AAEL012211 PE=4 SV=1
CCG024178.1	1245	2	403	0.06902769	11.31581168	7.356949232	Up	9.36E-05	0.00015346	AAEL015402-FA OS=Aedes aegypti GN=AAEL015402 PE=4 SV=1
CCG005017.1	810	69	403	3.6603848	17.39282165	2.248424776	Up	0	0	AAEL014038-FA OS=Aedes aegypti GN=AAEL014038 PE=3 SV=1
CCG027722.1	10348	849	403	3.52544498	1.361440427	-1.372671511	Down	3.74E-60	1.80E-59	AAEL006808-FA OS=Aedes aegypti GN=AAEL006808 PE=4 SV=1
CCG007777.1	4587	31	404	0.29039934	3.078950036	3.406328376	Up	2.40E-13	6.76E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010464 PE=4 SV=1
CCG016418.2	2184	60	404	1.18048721	6.466640941	2.453634091	Up	0	0	AAEL004136-FA OS=Aedes aegypti GN=AAEL004136 PE=4 SV=1
CCG014375.1	1339	74	404	2.37472768	10.54753086	2.151071321	Up	0	0	AAEL009087-FA OS=Aedes aegypti GN=AAEL009087 PE=4 SV=1
CCG001317.1	4845	122	404	1.08200364	2.914993564	1.429787349	Up	8.04E-14	2.58E-13	AAEL001089-FA OS=Aedes aegypti GN=AAEL001089 PE=4 SV=1
CCG027281.1	1061	146	404	5.91289468	13.31116288	1.170700128	Up	9.33E-15	3.34E-14	AAEL007885-FA OS=Aedes aegypti GN=AAEL007885 PE=4 SV=1
CCG010993.1	1125	100	405	3.81953197	12.58497964	1.720235112	Up	0	0	AAEL008459-FA OS=Aedes aegypti GN=AAEL008459 PE=4 SV=1
CCG000775.1	1301	114	405	3.76521887	10.88247663	1.531201287	Up	8.48E-14	2.70E-13	Quinone oxidoreductase OS=Culex quinquefasciatus GN=CpipJ_CPIJ019133 PE=4 SV=1
CCG009341.1	1357	138	405	4.36980352	10.433384	1.255566845	Up	1.49E-13	4.45E-13	AAEL003092-FA OS=Aedes aegypti GN=AAEL003092 PE=4 SV=1
CCG007553.1	1914	1285	405	28.8485418	7.39712753	-1.963461342	Down	4.53E-150	2.57E-149	-

CCG016870.1	1194	26	406	0.93568936	11.88698523	3.667209403	Up	4.60E-12	1.05E-11	AAEL005326-PA OS=Aedes aegypti GN=AAEL005326 PE=4 SV=1
CCG000406.1	2657	99	406	1.60105522	5.341761524	1.738292501	Up	0	0	AAEL004545-PA (Fragment) OS=Aedes aegypti GN=AAEL004545 PE=4 SV=1
CCG006302.1	471	141	406	12.8635511	30.13388613	1.228097769	Up	0	0	AAEL004726-PA OS=Aedes aegypti GN=AAEL004726 PE=4 SV=1
CCG010438.1	522	162	406	13.3354349	27.18977082	1.027799118	Up	0	0	AAEL007589-PA OS=Aedes aegypti GN=AAEL007589 PE=4 SV=1
CCG028151.1	5484	1092	406	8.55633665	2.588085407	-1.72510802	Down	3.90E-107	2.08E-106	AAEL012889-PA (Fragment) OS=Aedes aegypti GN=AAEL012889 PE=4 SV=1
CCG003413.1	1599	31	407	0.83305927	8.89807295	3.417001878	Up	2.40E-13	6.76E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003716 PE=4 SV=1
CCG007411.1	627	79	407	5.4140495	22.69221475	2.06741744	Up	0	0	AAEL004859-PA (Fragment) OS=Aedes aegypti GN=AAEL004859 PE=4 SV=1
CCG015246.1	870	128	407	6.32198394	16.35404442	1.371198188	Up	4.15E-14	1.42E-13	AAEL002686-PA OS=Aedes aegypti GN=AAEL002686 PE=3 SV=1
CCG028244.1	1467	162	407	4.74512407	9.69871755	1.031348185	Up	0	0	AGAP006038-PA OS=Anopheles gambiae GN=AGAP006038 PE=4 SV=4
CCG007005.1	2313	1289	407	23.9463847	6.151326695	-1.96083836	Down	2.90E-150	1.65E-149	AAEL014864-PA OS=Aedes aegypti GN=AAEL014864 PE=4 SV=1
CCG016753.1	1086	1473	407	58.2821539	13.1013063	-2.153343527	Down	1.07E-193	6.36E-193	-
CCG005178.3	2430	105	408	1.85671693	5.869537828	1.660493028	Up	0	0	Scarfoid attachment factor B OS=Culex quinquefasciatus GN=CpipJ_CPIJ017179 PE=1 SV=1
CCG000583.2	1677	116	408	2.97226548	8.505054814	1.516757551	Up	1.16E-13	3.59E-13	AAEL006508-PA OS=Aedes aegypti GN=AAEL006508 PE=4 SV=1
CCG003538.1	2631	2077	408	33.9217555	5.421123878	-2.645546955	Down	0	0	AAEL002070-PA (Fragment) OS=Aedes aegypti GN=AAEL002070 PE=4 SV=1

CCG021900.1	939	47	409	2.15077479	15.22676805	2.823681385	Up	0	0	AAEL008691-PA OS=Aedes aegypti GN=AAEL008691 PE=4 SV=1
CCG018751.1	1773	51	409	1.23601605	8.064261252	2.705844895	Up	0	0	AAEL008658-PA OS=Aedes aegypti GN=AAEL008658 PE=4 SV=1
CCG012418.1	1239	107	409	3.71086489	11.53989927	1.63680325	Up	0	0	AAEL007239-PA OS=Aedes aegypti GN=AAEL007239 PE=4 SV=1
CCG025000.1	670	853	409	54.7062442	21.34020179	-1.358131695	Down	1.79E-59	8.61E-59	AAEL006918-PB OS=Aedes aegypti GN=AAEL006918 PE=4 SV=1
CCG005323.1	2689	11731	409	187.459262	5.317194199	-5.139768143	Down	0	0	AGAP000113-PA (Fragment) OS=Anopheles gambiae GN=AgaP_AGAP000113 PE=4 SV=5
CCG005336.1	1040	54	410	2.23112084	13.78162834	2.626905801	Up	0	0	Dihydrodipicolinate reductase OS=Culex quinquefasciatus GN=CpipJ_CPIJ002390 PE=4 SV=1
CCG016692.1	2100	96	410	1.96433073	6.82518737	1.796830803	Up	0	0	AAEL005634-PA OS=Aedes aegypti GN=AAEL005634 PE=4 SV=1
CCG024392.1	522	117	410	9.63114741	27.45765034	1.511428584	Up	9.44E-14	2.97E-13	AAEL003401-PA OS=Aedes aegypti GN=AAEL003401 PE=4 SV=1
CCG008416.2	1047	5621	410	230.690428	13.68948756	-4.074817796	Down	0	0	AAEL003067-PB OS=Aedes aegypti GN=AAEL003067 PE=4 SV=1
CCG017379.1	591	29	411	2.10849798	24.31108588	3.527326792	Up	6.66E-13	1.69E-12	AAEL010150-PA OS=Aedes aegypti GN=AAEL010150 PE=4 SV=1
CCG028530.1	1295	31	411	1.02861913	11.09486622	3.431111477	Up	2.40E-13	6.77E-13	Atfermis protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014048 PE=4 SV=1
CCG017311.1	2355	36	411	0.65686219	6.100998621	3.215382786	Up	6.13E-14	2.03E-13	AAEL002322-PA OS=Aedes aegypti GN=AAEL002322 PE=4 SV=1
CCG016785.1	1905	49	411	1.10525827	7.542179398	2.770597943	Up	0	0	AAEL011664-PA OS=Aedes aegypti GN=AAEL011664 PE=4 SV=1
CCG016932.1	4641	95	411	0.87957871	3.095852565	1.815452179	Up	0	0	ATYP4-binding cassette sub-family A member 3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004978 PE=3 SV=1

CCG007948.1	10466	1782	411	7.31626859	1.37281213	-2.413973834	Down	2.07E-269	1.29E-268	AAEL005903-PA OS=Aedes aegypti GN=AAEL005903 PE=4 SV=1
CCG016751.1	536	25	412	2.00418538	26.87091424	3.744957541	Up	9.32E-12	2.08E-11	AAEL004842-PA OS=Aedes aegypti GN=AAEL004842 PE=4 SV=1
CCG013260.1	534	44	412	3.54057738	26.97155436	2.929382112	Up	3.91E-14	1.34E-13	AAEL009759-PA OS=Aedes aegypti GN=AAEL009759 PE=4 SV=1
CCG018925.1	14914	94	412	0.27082976	0.965724154	1.834224879	Up	0	0	AAEL011396-PA OS=Aedes aegypti GN=AAEL011396 PE=4 SV=1
CCG026852.1	1185	129	412	4.67771795	12.15427007	1.377586476	Up	0	0	Uricase OS=Aedes aegypti GN=UO PE=2 SV=1
CCG016856.2	1962	134	412	2.93473213	7.340881769	1.322724541	Up	0	0	AAEL014112-PA OS=Aedes aegypti GN=AAEL014112 PE=4 SV=1
CCG003642.1	537	150	412	12.0027192	26.82087529	1.159995041	Up	0	0	AAEL006771-PA OS=Aedes aegypti GN=AAEL006771 PE=4 SV=1
CCG009784.1	1325	152	412	4.92935823	10.87004531	1.140886218	Up	8.39E-14	2.69E-13	AAEL000562-PA OS=Aedes aegypti GN=AAEL000562 PE=4 SV=1
CCG010001.1	1815	932	412	22.0648995	7.935432524	-1.475372414	Down	2.68E-73	1.34E-72	AAEL009038-PA (Fragment) OS=Aedes aegypti GN=AAEL009038 PE=4 SV=1
CCG021923.1	1777	1084	412	26.2122635	8.105126635	-1.69333531	Down	1.26E-103	6.73E-103	AAEL000271-PA OS=Aedes aegypti GN=AAEL000271 PE=4 SV=1
CCG018684.1	372	2467	412	284.963267	38.71723126	-2.879728269	Down	0	0	AAEL004829-PA OS=Aedes aegypti GN=AAEL004829 PE=4 SV=1
CCG002380.1	1260	15	413	0.51154446	11.45854628	4.48542058	Up	1.48E-08	2.93E-08	AAEL003004-PA OS=Aedes aegypti GN=AAEL003004 PE=4 SV=1
CCG011975.1	1719	55	414	1.37483153	8.419270846	2.614440448	Up	0	0	AAEL009187-PA OS=Aedes aegypti GN=AAEL009187 PE=4 SV=1
CCG015700.1	2952	109	414	1.5866196	4.902685157	1.627615837	Up	0	0	AAEL004387-PA OS=Aedes aegypti GN=AAEL004387 PE=4 SV=1
CCG016864.1	1011	125	414	5.31277629	14.31525874	1.430015877	Up	2.35E-14	8.22E-14	AAEL007105-PA OS=Aedes aegypti GN=AAEL007105 PE=4 SV=1
CCG023434.1	1896	23	415	0.52125733	7.651732521	3.875718774	Up	4.01E-11	8.67E-11	AAEL009397-PA OS=Aedes aegypti GN=AAEL009397 PE=4 SV=1
CCG021092.1	717	39	415	2.3372659	20.2338701	3.113878511	Up	7.33E-14	2.38E-13	AAEL008400-PA OS=Aedes aegypti GN=AAEL008400 PE=4 SV=1

CCG012728.1	2326	54	415	0.99757767	6.237181797	2.644393228	Up	0	0	AAEL008865-PA OS=Aedes aegypti GN=AAEL008865 PE=3 SV=1
CCG024253.1	1494	82	415	2.35844594	9.710632437	2.041728725	Up	0	0	Uncharacterized protein OS=Danio rerio GN=BX005085.3 PE=1 SV=1
CCG003409.1	1242	133	415	4.60142891	11.68090568	1.343998295	Up	9.90E-14	3.11E-13	AAEL004420-PA OS=Aedes aegypti GN=AAEL004420 PE=4 SV=1
CCG002505.1	2571	135	415	2.25628711	5.642817915	1.322465133	Up	1.90E-13	5.49E-13	AAEL002496-PA OS=Aedes aegypti GN=AAEL002496 PE=4 SV=1
CCG019136.1	668	1127	415	72.4953457	21.71809111	-1.73899107	Down	9.39E-112	5.06E-111	AAEL002293-PA OS=Aedes aegypti GN=AAEL002293 PE=4 SV=1
CCG012169.1	426	115	416	11.599811	34.13765995	1.557262871	Up	6.13E-14	2.02E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005871 PE=4 SV=1
CCG017372.1	732	120	416	7.04421879	19.86699882	1.495862326	Up	0	0	AAEL010419-PA OS=Aedes aegypti GN=AAEL010419 PE=4 SV=1
CCG026897.2	1221	132	416	4.64537671	11.91043664	1.358358803	Up	1.46E-13	4.36E-13	AAEL001448-PA OS=Aedes aegypti GN=AAEL001448 PE=4 SV=1
CCG021064.1	603	838	416	59.7158169	24.1171528	-1.308053512	Down	3.02E-55	1.44E-54	AAEL008054-PA OS=Aedes aegypti GN=AAEL008054 PE=4 SV=1
CCG011764.1	5541	4398	416	34.1059182	2.624552091	-3.699878967	Down	0	0	AAEL011436-PA OS=Aedes aegypti GN=AAEL011436 PE=4 SV=1
CCG027199.1	458	0	417	0.001	31.82882405	14.95804624	Up	4.44E-16	1.65E-15	Histone H2A OS=Aedes aegypti GN=AAEL003669 PE=3 SV=1
CCG000563.1	1086	27	417	1.06830832	13.42320572	3.651329275	Up	2.28E-12	5.41E-12	AAEL013795-PA OS=Aedes aegypti GN=AAEL013795 PE=4 SV=1
CCG017369.1	2133	38	417	0.76551801	6.834318525	3.158289264	Up	4.26E-14	1.45E-13	AAEL007211-PA OS=Aedes aegypti GN=AAEL007211 PE=4 SV=1
CCG024267.1	717	48	417	2.87663495	20.33138273	2.821254277	Up	5.35E-14	1.79E-13	AAEL008038-PA OS=Aedes aegypti GN=AAEL008038 PE=4 SV=1
CCG002386.1	1358	73	417	2.30986055	10.73461076	2.216392218	Up	0	0	AAEL010782-PA OS=Aedes aegypti GN=AAEL010782 PE=3 SV=1
CCG010331.1	3849	1714	417	19.1348727	3.787373711	-2.336934617	Down	2.99E-249	1.84E-248	AAEL002583-PA OS=Aedes aegypti GN=TOLL7 PE=4 SV=1
CCG008256.1	1332	4	418	0.12903824	10.97039016	6.409672336	Up	2.95E-05	5.04E-05	AAEL010627-PA OS=Aedes aegypti GN=AAEL010627 PE=4 SV=1

CCG013026.1	1089	90	418	3.55121774	13.41832846	1.91781924	Up	0	0	AAEL014272-PA OS=Aedes aegypti GN=AAEL014272 PE=4 SV=1
CCG017651.1	3423	117	418	1.46872888	4.268933594	1.539307616	Up	9.44E-14	2.98E-13	AAEL011978-PA OS=Aedes aegypti GN=AAEL011978 PE=4 SV=1
CCG024355.1	594	160	418	11.5743393	24.60026884	1.087744241	Up	4.24E-14	1.44E-13	AAEL005313-PA OS=Aedes aegypti GN=AAEL005313 PE=4 SV=1
CCG004542.1	4572	2317	418	21.7762194	3.19609792	-2.768369993	Down	0	0	AAEL008421-PA OS=Aedes aegypti GN=AAEL008421 PE=4 SV=1
CCG027229.1	742	2686	418	155.548123	19.69347667	-2.981571254	Down	0	0	AAEL014944-PA OS=Aedes aegypti GN=SGP2D7 PE=2 SV=1
CCG023633.1	1392	15	419	0.46303593	10.52264222	4.506229042	Up	1.48E-08	2.93E-08	AAEL007939-PA OS=Aedes aegypti GN=AAEL007939 PE=4 SV=1
CCG014896.1	411	146	419	15.264188	35.63872985	1.223295079	Up	9.33E-15	3.34E-14	Ribosomal protein OS=Aedes aegypti GN=AAEL009613 PE=3 SV=1
CCG019497.1	2685	167	419	2.67260547	5.455313954	1.029415345	Up	1.62E-13	4.78E-13	AAEL009404-PA OS=Aedes aegypti GN=AAEL009404 PE=4 SV=1
CCG003795.1	492	776	419	67.7734026	29.77137798	-1.186793205	Down	3.19E-44	1.45E-43	AAEL001952-PA OS=Aedes aegypti GN=AAEL001952 PE=2 SV=1
CCG022747.1	2253	1002	419	19.1103746	6.501339533	-1.555547156	Down	5.55E-85	2.85E-84	AAEL001207-PA OS=Aedes aegypti GN=AAEL001207 PE=4 SV=2
CCG009744.1	1365	62	420	1.95173886	10.75639285	2.462362411	Up	0	0	AAEL005374-PA OS=Aedes aegypti GN=AAEL003374 PE=4 SV=1
CCG009279.1	1179	107	420	3.89971298	12.45333015	1.675091735	Up	0	0	AAEL006286-PA OS=Aedes aegypti GN=AAEL006286 PE=4 SV=1
CCG014462.1	270	129	420	20.5299843	54.37954165	1.405331466	Up	0	0	Activating transcription factor OS=Aedes aegypti PE=2 SV=1
CCG003181.1	1761	58	421	1.41524396	8.357430165	2.562008632	Up	0	0	AAEL014483-PA OS=Aedes aegypti GN=AAEL014483 PE=4 SV=1
CCG000387.1	1254	92	421	3.15248452	11.73639117	1.896427671	Up	1.44E-14	5.12E-14	AAEL010376-PA OS=Aedes aegypti GN=AAEL010376 PE=4 SV=1
CCG003424.1	1098	126	421	4.93095315	13.40385658	1.442709703	Up	1.34E-13	4.06E-13	AAEL008850-PA OS=Aedes aegypti GN=AAEL008850 PE=4 SV=1
CCG024082.1	1458	746	421	21.9858862	10.09426236	-1.123042193	Down	4.23E-39	1.88E-38	AAEL006542-PA OS=Aedes aegypti GN=AAEL006542 PE=4 SV=1

CCG000658.1	1230	1062	421	37.1006977	11.96539392	-1.632578424	Down	1.96E-96	1.03E-95	Possible mucin OS=Aedes aegypti PE=2 SV=1
CCG022816.1	3568	3868	421	46.5826607	4.124841514	-3.497382453	Down	0	0	AAEL003135-PA OS=Aedes aegypti GN=AAEL003135 PE=4 SV=1
CCG016906.1	1424	6700	421	202.175015	10.33527705	-4.289955753	Down	0	0	AAEL015304-PA OS=Aedes aegypti GN=AAEL015304 PE=4 SV=1
CCG025737.1	441	0	422	0.001	33.45213786	15.02981079	Up	4.44E-16	1.64E-15	-
CCG017096.2	684	32	422	2.01027998	21.56782573	3.423412393	Up	1.77E-13	5.17E-13	AAEL003332-PA OS=Aedes aegypti GN=AAEL003332 PE=4 SV=1
CCG019231.1	1679	94	422	2.40569092	8.786416199	1.868823541	Up	0	0	AAEL001778-PA OS=Aedes aegypti GN=AAEL001778 PE=4 SV=1
CCG007829.1	1530	107	422	3.00507294	9.642086796	1.681945406	Up	0	0	AAEL005053-PA OS=Aedes aegypti GN=AAEL005053 PE=4 SV=1
CCG021650.1	2472	135	422	2.34664813	5.967796439	1.346596795	Up	1.90E-13	5.49E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000842 PE=4 SV=1
CCG026949.1	973	146	422	6.4476683	15.16176033	1.233587834	Up	9.33E-15	3.33E-14	AAEL010250-PA OS=Aedes aegypti GN=AAEL010250 PE=4 SV=1
CCG021744.1	1692	34	423	0.86345802	8.739569193	3.339364216	Up	8.82E-14	2.79E-13	AAEL008116-PA OS=Aedes aegypti GN=AAEL008116 PE=4 SV=1
CCG008126.1	2547	64	423	1.07972635	5.805791549	2.426827057	Up	0	0	AAEL004200-PA OS=Aedes aegypti GN=AAEL004200 PE=3 SV=1
CCG025895.2	959	96	423	4.30145414	15.41955274	1.841864556	Up	0	0	-
CCG006202.1	978	146	423	6.41470476	15.11999087	1.237002498	Up	9.33E-15	3.34E-14	AAEL006067-PA OS=Aedes aegypti GN=AAEL006067 PE=4 SV=1
CCG011114.1	621	17	424	1.17630513	23.86845306	4.342770817	Up	3.39E-09	6.89E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014666 PE=4 SV=1
CCG013983.1	3288	41	424	0.53581482	4.508001628	3.072681654	Up	1.02E-13	3.19E-13	AAEL004243-PA OS=Aedes aegypti GN=AAEL004243 PE=4 SV=1
CCG025303.1	822	42	424	2.19553389	18.03200651	3.037916236	Up	7.66E-14	2.48E-13	Clp-domain serine protease OS=Culex quinquefasciatus GN=CpipJ_CPIJ000615 PE=3 SV=1
CCG013290.1	1547	47	424	1.30547998	9.581324726	2.875644807	Up	0	0	AAEL011362-PA OS=Aedes aegypti GN=AAEL011362 PE=4 SV=1

CCG011243.1	417	54	424	5.56442606	35.54510636	2.675346156	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007523 PE=4 SV=1
CCG025412.1	1560	92	424	2.53411255	9.501480353	1.906671702	Up	1.44E-14	5.12E-14	AAEL009240-PA OS=Aedes aegypti GN=AAEL009246 PE=3 SV=1
CCG018233.1	769	113	424	6.31414826	19.27478459	1.610054696	Up	1.42E-13	4.26E-13	AAEL008714-PA OS=Aedes aegypti GN=AAEL008714 PE=4 SV=1
CCG011168.2	2364	148	424	2.69015259	6.270012416	1.220780293	Up	0	0	Gastrin zinc finger protein XLCGF8.2DB OS=Culex quinquefasciatus GN=CpipJ_CPIJ012331 PE=4 SV=1
CCG013615.1	969	1608	424	71.3058135	15.29650088	-2.220818033	Down	1.13E-219	6.86E-219	AAEL013656-PA OS=Aedes aegypti GN=AAEL013656 PE=4 SV=1
CCG009808.1	3606	1894	424	22.5692394	4.110457391	-2.456986957	Down	2.69E-292	1.69E-291	AAEL005437-PA OS=Aedes aegypti GN=AAEL005437 PE=4 SV=1
CCG008322.1	1770	4713	424	114.416022	8.374186074	-3.772196308	Down	0	0	AAEL013526-PA OS=Aedes aegypti GN=AAEL013526 PE=4 SV=1
CCG001171.1	1314	21	425	0.68673092	11.30690078	4.041314812	Up	1.76E-10	3.72E-10	AAEL009401-PA OS=Aedes aegypti GN=AAEL009401 PE=4 SV=1
CCG007794.1	2385	70	425	1.26116622	6.229462318	2.304349218	Up	0	0	AAEL014631-PA OS=Aedes aegypti GN=AAEL014631 PE=4 SV=1
CCG025982.1	1996	85	425	1.82987347	7.443520856	2.024241299	Up	0	0	AAEL001996-PA OS=Aedes aegypti GN=AAEL001996 PE=4 SV=1
CCG005360.1	2595	707	425	11.7069759	5.72534398	-1.03193417	Down	1.48E-32	6.33E-32	AAEL010804-PA OS=Aedes aegypti GN=AAEL010804 PE=4 SV=1
CCG007479.1	415	895	425	92.6696686	35.80064489	-1.372111637	Down	2.82E-63	1.37E-62	AAEL002526-PA OS=Aedes aegypti GN=AAEL002526 PE=4 SV=1
CCG028061.1	2110	12	426	0.24437764	7.057926969	4.852060323	Up	1.30E-07	2.49E-07	AAEL000918-PA (Fragment) OS=Aedes aegypti GN=AAEL000918 PE=4 SV=2
CCG007771.1	759	136	426	7.69945179	19.62085099	1.349559983	Up	1.70E-13	4.99E-13	AAEL009497-PA OS=Aedes aegypti GN=AAEL009497 PE=4 SV=1
CCG005113.1	2079	146	426	3.0175956	7.163167823	1.247198265	Up	9.33E-15	3.34E-14	AAEL001163-PA OS=Aedes aegypti GN=AAEL001163 PE=4 SV=1

CCG018453.1	1475	157	426	4.57372768	10.09642434	1.142402075	Up	0	0	Delta-aminolevulinic acid dehydratase OS=Aedes aegypti GN=AAEL002304 PE=2 SV=1
CCG000593.1	954	755	426	34.0064462	15.61029969	-1.12331001	Down	1.45E-39	6.45E-39	homeobox protein extradenticle OS=Culex quinquefasciatus GN=CpipJ_CPIJ008039 PE=3 SV=1
CCG025568.1	2583	1016	426	16.9017617	5.765476541	-1.551661863	Down	7.85E-86	4.04E-85	AAEL011280-PA (Fragment) OS=Aedes aegypti GN=AAEL011280 PE=3 SV=1
CCG028339.1	2370	11	427	0.19943759	6.29838995	4.980973845	Up	2.67E-07	5.03E-07	AAEL000730-PA OS=Aedes aegypti GN=AAEL000730 PE=4 SV=1
CCG000926.1	1473	27	427	0.78763261	10.1338657	3.685517961	Up	2.28E-12	5.40E-12	AAEL013312-PA (Fragment) OS=Aedes aegypti GN=AAEL013312 PE=4 SV=1
CCG003193.1	2004	58	427	1.24363503	7.448694701	2.582424468	Up	0	0	zinc finger protein 394 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002919 PE=4 SV=1
CCG009635.1	1819	92	427	2.17329059	8.206258484	1.916843507	Up	1.44E-14	5.12E-14	AAEL001561-PA OS=Aedes aegypti GN=AAEL001561 PE=4 SV=1
CCG012584.1	256	0	428	0.001	58.44586898	15.83481344	Up	4.44E-16	1.65E-15	histone H2B OS=Ixodes scapularis GN=IscW_ISCW019496 PE=2 SV=1
CCG005390.1	1138	25	428	0.94397484	13.1477526	3.799924	Up	9.32E-12	2.08E-11	AAEL006407-PA OS=Aedes aegypti GN=AAEL006407 PE=4 SV=1
CCG027125.1	1471	63	428	1.84030814	10.17140888	2.466500267	Up	0	0	Queuine tRNA-ribosyltransferase OS=Aedes aegypti GN=AAEL010993 PE=3 SV=1
CCG006531.1	457	70	428	6.58179742	32.73991785	2.314497173	Up	0	0	AAEL003029-PA OS=Aedes aegypti GN=AAEL003029 PE=4 SV=1
CCG010601.1	1167	76	428	2.79837175	12.82103038	2.195852677	Up	0	0	AAEL001811-PA OS=Aedes aegypti GN=AAEL001811 PE=4 SV=1
CCG025979.1	884	111	428	5.39552097	16.92550052	1.649364324	Up	0	0	AAEL001997-PA OS=Aedes aegypti GN=AAEL001997 PE=4 SV=1
CCG016459.1	1443	169	428	5.03249144	10.36877509	1.042900754	Up	0	0	AAEL004247-PA OS=Aedes aegypti GN=AAEL004247 PE=4 SV=1

CCG013087.1	1170	18	429	0.66107284	12.81803482	4.27722204	Up	1.62E-09	3.33E-09	AAEL004318-PA OS=Aedes aegypti GN=AAEL004318 PE=4 SV=1
CCG011005.1	1566	27	429	0.74085749	9.576692679	3.692259539	Up	2.28E-12	5.42E-12	AAEL010185-PA OS=Aedes aegypti GN=AAEL010185 PE=4 SV=1
CCG023370.1	1120	51	429	1.95665756	13.39026851	2.774721699	Up	0	0	AAEL000697-PA OS=Aedes aegypti GN=AAEL006971 PE=4 SV=1
CCG009833.1	3887	72	429	0.79594054	3.858271349	2.27722204	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014679 PE=4 SV=1
CCG013217.1	1203	144	429	5.14350938	12.4664179	1.27722204	Up	7.53E-14	2.43E-13	AAEL003750-PA OS=Aedes aegypti GN=AAEL003750 PE=4 SV=1
CCG013426.1	729	2	430	0.11788679	20.62010838	7.450506053	Up	9.36E-05	0.00015377	AAEL014270-PA OS=Aedes aegypti GN=AAEL014276 PE=4 SV=1
CCG016381.1	1290	16	430	0.53295795	11.65275892	4.450506053	Up	7.09E-09	1.42E-08	DNA primase OS=Aedes aegypti GN=AAEL014314 PE=3 SV=1
CCG003700.1	1540	96	430	2.67863281	9.761077281	1.865543553	Up	0	0	AAEL005050-PA OS=Aedes aegypti GN=AAEL005050 PE=4 SV=1
CCG017775.1	1430	111	430	3.33541297	10.51192938	1.656090187	Up	0	0	AAEL012913-PA OS=Aedes aegypti GN=AAEL012913 PE=4 SV=1
CCG011041.1	3786	1948	430	22.1090975	3.970432914	-2.477271909	Down	1.60E-303	1.01E-302	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006592 PE=4 SV=1
CCG025371.1	540	21	431	1.67104523	27.90188387	4.06153984	Up	1.76E-10	3.73E-10	AAEL006610-PA OS=Aedes aegypti GN=AAEL006610 PE=4 SV=1
CCG016032.1	1347	23	431	0.73370742	11.18561046	3.930295307	Up	4.01E-11	8.66E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_17290 PE=4 SV=1
CCG009025.1	927	85	431	3.94005118	16.25352458	2.044466327	Up	0	0	AAEL012771-PA OS=Aedes aegypti GN=AAEL012771 PE=4 SV=1
CCG019356.1	504	89	431	7.58790948	29.89487557	1.978123832	Up	0	0	AAEL013778-PA OS=Aedes aegypti GN=AAEL013778 PE=4 SV=1
CCG013509.1	1686	99	431	2.52313388	8.936546435	1.824500643	Up	0	0	AAEL014321-PA OS=Aedes aegypti GN=AAEL014321 PE=4 SV=1

CCG016987.1	306	128	431	17.9742681	49.23861859	1.453857263	Up	4.15E-14	1.42E-13	Putative uncharacterized protein OS=Ochlerotatus taeniorhynchus PE=4 SV=1
CCG015999.1	2117	132	431	2.67926546	7.117155073	1.409463144	Up	1.46E-13	4.35E-13	AAEL012975-PA OS=Aedes aegypti GN=AAEL012975 PE=4 SV=1
CCG024279.1	1631	167	431	4.39972145	9.237901465	1.07015297	Up	1.62E-13	4.78E-13	AAEL007343-PA OS=Aedes aegypti GN=AAEL007343 PE=4 SV=1
CCG017769.1	1905	1407	431	31.7367016	7.909195427	-2.00454935	Down	1.06E-168	6.17E-168	AAEL006216-PA OS=Aedes aegypti GN=AAEL006216 PE=4 SV=1
CCG019138.2	663	2422	431	156.972394	22.72551627	-2.788125887	Down	0	0	AAEL002309-PA OS=Aedes aegypti GN=TPX4 PE=4 SV=1
CCG016602.1	3285	2630	431	34.4019489	4.586611047	-2.906989821	Down	0	0	AAEL008032-PA OS=Aedes aegypti GN=AAEL008032 PE=4 SV=1
CCG007919.1	2084	49	432	1.01032485	7.24662935	2.842490862	Up	0	0	AAEL006858-PA OS=Aedes aegypti GN=AAEL006858 PE=4 SV=1
CCG004873.1	1391	86	432	2.65664786	10.85691989	2.030935951	Up	0	0	AAEL007652-PA OS=Aedes aegypti GN=AAEL007652 PE=4 SV=1
CCG021884.1	771	107	432	5.96337432	19.58751695	1.71573372	Up	0	0	AAEL012238-PA OS=Aedes aegypti GN=AAEL012238 PE=2 SV=1
CCG020004.1	586	14	433	1.02658069	25.83094512	4.653181496	Up	3.07E-08	6.01E-08	-
CCG026194.1	930	18	433	0.83167228	16.27627295	4.290611417	Up	1.62E-09	3.33E-09	-
CCG005507.2	1518	75	433	2.1230106	9.971629672	2.231717728	Up	0	0	AAEL001395-PA OS=Aedes aegypti GN=AAEL001395 PE=4 SV=1
CCG021645.1	1258	86	433	2.93751763	12.03253883	2.034271664	Up	0	0	-
CCG010065.1	492	88	433	7.68564359	30.7661257	2.0011048	Up	2.93E-14	1.02E-13	AAEL006502-PA OS=Aedes aegypti GN=AAEL006502 PE=4 SV=1
CCG001897.1	1182	107	433	3.88981523	12.8062046	1.719069432	Up	0	0	AAEL010388-PA OS=Aedes aegypti GN=AAEL010388 PE=4 SV=1
CCG025038.1	1583	111	433	3.01303888	9.562181833	1.666120552	Up	0	0	AAEL005395-PA OS=Aedes aegypti GN=AAEL005395 PE=4 SV=1
CCG004514.1	1582	761	433	20.6700177	9.568226196	-1.111216225	Down	3.35E-39	1.49E-38	AGAP007599-PA OS=Anopheles gambiae GN=AGAP007599 PE=4 SV=4

CCG003186.1	360	1417	433	169.13365	42.04703845	-2.008087624	Down	2.80E-170	1.62E-169	AAEL005667-PA (Fragment) OS=Aedes aegypti GN=AAEL005667 PE=4 SV=1
CCG014520.1	1077	23	434	0.91764521	14.08717931	3.94030248	Up	4.01E-11	8.67E-11	AAEL007581-PA OS=Aedes aegypti GN=AAEL007581 PE=4 SV=1
CCG005560.1	2110	32	434	0.6516737	7.190470199	3.463864436	Up	1.77E-13	5.17E-13	AAEL003549-PA OS=Aedes aegypti GN=AAEL003549 PE=4 SV=1
CCG023979.1	1389	44	434	1.3611723	10.92288849	3.004432818	Up	3.91E-14	1.34E-13	AAEL000289-PA OS=Aedes aegypti GN=AAEL000289 PE=4 SV=1
CCG015551.1	3743	78	434	0.89544197	4.053404253	2.178462217	Up	0	0	AAEL003036-PA OS=Aedes aegypti GN=AAEL003036 PE=4 SV=1
CCG005556.1	636	93	434	6.28331025	23.85517629	1.924705625	Up	4.04E-14	1.39E-13	-
CCG007129.4	1686	128	434	3.26223371	8.998749774	1.463864436	Up	4.15E-14	1.42E-13	AAEL005820-PA OS=Aedes aegypti GN=AAEL005820 PE=4 SV=1
CCG014872.1	2187	175	434	3.43836468	6.937307782	1.012653324	Up	3.55E-13	9.58E-13	AAEL008533-PA OS=Aedes aegypti GN=AAEL008533 PE=4 SV=1
CCG009863.1	1725	37	435	0.92166967	8.815565447	3.257731429	Up	5.17E-14	1.74E-13	AAEL012465-PA OS=Aedes aegypti GN=AAEL012465 PE=4 SV=1
CCG006096.1	3450	1010	435	12.5795455	4.407782723	-1.512954783	Down	2.95E-82	1.51E-81	AAEL004690-PA OS=Aedes aegypti GN=AAEL004690 PE=4 SV=1
CCG020898.1	1158	26	436	0.96477815	13.16218366	3.77005781	Up	4.60E-12	1.05E-11	AAEL013531-PA OS=Aedes aegypti GN=AAEL013531 PE=3 SV=1
CCG017303.1	942	98	436	4.47031209	16.18026398	1.855787684	Up	0	0	AAEL002362-PA OS=Aedes aegypti GN=AAEL002362 PE=4 SV=1
CCG022027.1	1330	113	436	3.65081204	11.46000652	1.650318566	Up	1.42E-13	4.27E-13	AAEL014961-PA OS=Aedes aegypti GN=AAEL014961 PE=4 SV=1
CCG003576.1	1611	122	436	3.25407053	9.461085458	1.539760191	Up	8.04E-14	2.58E-13	-
CCG011407.1	1329	135	436	4.36487146	11.46862955	1.393681932	Up	1.90E-13	5.49E-13	AAEL009635-PA OS=Aedes aegypti GN=AAEL009635 PE=4 SV=1
CCG015962.1	1083	161	436	6.38792915	14.07369222	1.13958065	Up	0	0	Conserved membrane protein OS=Aedes aegypti PE=2 SV=1
CCG023643.1	1411	3240	436	98.6689866	10.8021323	-3.191280569	Down	0	0	AAEL003611-PA OS=Aedes aegypti GN=AAEL003611 PE=3 SV=1
CCG010646.1	477	7	437	0.63058311	32.02676509	5.666447751	Up	4.31E-06	7.69E-06	Histone H2A OS=Gasterosteus aculeatus PE=3 SV=1

CCG001168.1	1296	81	437	2.68560841	11.78762882	2.13395267	Up	4.46E-14	1.51E-13	AAEL007926-PA OS=Aedes aegypti GN=AAEL007926 PE=4 SV=1
CCG005773.1	1390	104	437	3.21500173	10.99047982	1.773362955	Up	0	0	AAEL014705-PA OS=Aedes aegypti GN=AAEL014705 PE=4 SV=1
CCG010154.1	967	129	437	5.73226036	15.79810439	1.462575418	Up	0	0	-
CCG012396.1	1986	174	437	3.76471995	7.692229078	1.030859177	Up	1.78E-13	5.20E-13	AAEL011277-PA (Fragment) OS=Aedes aegypti GN=AAEL011277 PE=4 SV=1
CCG016523.1	920	1058	437	49.4151948	16.60518147	-1.573321239	Down	4.11E-91	2.14E-90	AAEL003482-PA OS=Aedes aegypti GN=AAEL003482 PE=4 SV=1
CCG017921.1	2217	34	438	0.65898556	6.906506642	3.389637422	Up	8.82E-14	2.80E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014312 PE=4 SV=1
CCG024612.1	1509	79	438	2.24957524	10.14693521	2.173319515	Up	0	0	AAEL005468-PA OS=Aedes aegypti GN=AAEL005468 PE=4 SV=1
CCG019052.2	1428	719	438	21.6353216	10.72249666	-1.012747697	Down	4.06E-32	1.73E-31	AAEL000851-PA OS=Aedes aegypti GN=AAEL000851 PE=4 SV=1
CCG012053.1	2415	963	438	17.1345153	6.340258893	-1.434291724	Down	1.38E-72	6.88E-72	AAEL004122-PA OS=Aedes aegypti GN=AAEL004122 PE=4 SV=1
CCG020675.1	1229	18	439	0.62933704	12.4871306	4.310465332	Up	1.62E-09	3.32E-09	AAEL006787-PA OS=Aedes aegypti GN=AAEL006787 PE=4 SV=1
CCG027987.1	2469	29	439	0.50470729	6.215748685	3.622409338	Up	6.66E-13	1.69E-12	AAEL001591-PA OS=Aedes aegypti GN=AAEL001591 PE=4 SV=1
CCG006225.1	1061	108	439	4.37392209	14.46435768	1.725502831	Up	0	0	AAEL011782-PA OS=Aedes aegypti GN=AAEL011782 PE=4 SV=1
CCG017961.1	1605	131	439	3.50718706	9.561796575	1.446967332	Up	1.26E-13	3.87E-13	AAEL009701-PA (Fragment) OS=Aedes aegypti GN=AAEL009701 PE=4 SV=1
CCG000596.1	3465	1103	439	13.6783888	4.429057288	-1.626826742	Down	1.40E-99	7.41E-99	Uncharacterized protein OS=Anopheles darlingi GN=AND_13904 PE=4 SV=1
CCG021776.1	2286	3375	439	63.4395688	6.713334866	-3.240281453	Down	0	0	AAEL000153-PA OS=Aedes aegypti GN=AAEL000153 PE=4 SV=1
CCG023003.1	682	0	440	0.001	22.55372695	14.46107823	Up	4.44E-16	1.64E-15	AAEL008137-PA OS=Aedes aegypti GN=AAEL008137 PE=4 SV=1

CCG010136.1	1324	6	440	0.19472689	11.61755421	5.898710417	Up	8.38E-06	1.48E-05	Histone H2A OS=Gasterosteus aculeatus PE=3 SV=1
CCG009776.1	2265	1554	440	29.4812219	6.791011823	-2.118097871	Down	5.71E-200	3.41E-199	Alpha-amylase T OS=Culex quinquefasciatus GN=CpipJ_CPIJ017521 PE=3 SV=1
CCG020688.1	1572	6	441	0.16400662	9.806997491	5.901985549	Up	8.38E-06	1.48E-05	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001997 PE=4 SV=1
CCG003839.2	720	17	441	1.01456318	21.41194452	4.399485208	Up	3.39E-09	6.89E-09	AAEL001684-PA OS=Aedes aegypti GN=AAEL001684 PE=4 SV=1
CCG015950.3	1593	53	441	1.42962708	9.677715039	2.759027595	Up	0	0	AAEL005688-PA OS=Aedes aegypti GN=AAEL005688 PE=3 SV=1
CCG002811.1	477	60	441	5.40499806	32.31991626	2.580057454	Up	0	0	AAEL006006-PA OS=Aedes aegypti GN=AAEL006006 PE=4 SV=1
CCG023966.1	243	97	441	17.1525278	63.44279859	1.887035207	Up	3.62E-14	1.25E-13	AAEL008842-PA OS=Aedes aegypti GN=AAEL008842 PE=4 SV=1
CCG027956.1	445	106	441	10.2354873	34.64404507	1.759027595	Up	0	0	-
CCG008608.1	984	162	441	7.07428558	15.66727648	1.147098046	Up	0	0	AAEL010316-PA OS=Aedes aegypti GN=AAEL010316 PE=4 SV=1
CCG007729.1	999	998	441	42.9267219	15.43203209	-1.475947956	Down	2.10E-78	1.06E-77	AAEL008343-PA OS=Aedes aegypti GN=CYP4G35 PE=3 SV=1
CCG005307.1	1669	1347	441	34.6795881	9.237028194	-1.908586086	Down	2.32E-151	1.32E-150	Substrate carrier family 25 member 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004614 PE=4 SV=1
CCG016869.1	16943	1560	441	3.95637054	0.909909701	-2.120382264	Down	5.18E-201	3.09E-200	AAEL005337-PA OS=Aedes aegypti GN=AAEL005337 PE=4 SV=1
CCG025069.2	1005	9001	441	384.846349	15.33990055	-4.648921527	Down	0	0	AAEL006576-PA OS=Aedes aegypti GN=AAEL006576 PE=3 SV=1
CCG000536.1	1263	59	442	2.0072956	12.23401293	2.607572714	Up	0	0	AAEL013901-PA OS=Aedes aegypti GN=AAEL013901 PE=4 SV=1
CCG002720.1	3130	61	442	0.83742933	4.936600107	2.559478426	Up	1.75E-14	6.17E-14	Zinc finger protein 337 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014708 PE=4 SV=1
CCG028198.1	999	108	442	4.64537671	15.46702536	1.735328261	Up	0	0	Protein kinase C OS=Aedes aegypti GN=AAEL001108 PE=3 SV=1

CCG014716.1	2155	115	442	2.29304848	7.170096674	1.644725712	Up	6.13E-14	2.03E-13	AAEL001984-PA OS=Aedes aegypti GN=AAEL001984 PE=4 SV=1
CCG015677.1	2567	131	442	2.19284583	6.019305934	1.456792762	Up	1.26E-13	3.87E-13	AAEL001470-PA OS=Aedes aegypti GN=AAEL001470 PE=4 SV=1
CCG013256.1	18873	1068	442	2.43160476	0.818712358	-1.570480168	Down	1.02E-91	5.34E-91	Uncharacterized protein OS=Anopheles darlingi GN=AND_14085 PE=4 SV=1
CCG007046.2	1056	1865	442	75.8887832	14.63215751	-2.374744152	Down	2.30E-276	1.43E-275	Galactose-1-phosphate uridylyltransferase OS=Aedes aegypti GN=AAEL001607 PE=3 SV=1
CCG017640.1	1389	40	443	1.23742936	11.14940001	3.171547997	Up	6.86E-14	2.25E-13	AAEL000289-PA OS=Aedes aegypti GN=AAEL000289 PE=4 SV=1
CCG014888.1	903	114	443	5.42475055	17.15007377	1.660586078	Up	8.48E-14	2.70E-13	AAEL013177-PA OS=Aedes aegypti GN=AAEL013177 PE=4 SV=1
CCG015668.1	1199	127	443	4.5514231	12.916194	1.504791406	Up	1.57E-13	4.67E-13	AAEL007196-PA OS=Aedes aegypti GN=AAEL007196 PE=2 SV=1
CCG005749.1	873	163	443	8.02298596	17.73942338	1.144747938	Up	1.80E-13	5.24E-13	AAEL009703-PA OS=Aedes aegypti GN=AAEL009703 PE=4 SV=1
CCG008830.1	1476	1813	443	52.7805751	10.49221993	-2.330687117	Down	1.14E-262	7.05E-262	AAEL004440-PA OS=Aedes aegypti GN=AAEL004440 PE=4 SV=1
CCG006760.1	1870	2534	443	58.2274372	8.281559685	-2.813724717	Down	0	0	AAEL012200-PA OS=Aedes aegypti GN=CYP4C38 PE=3 SV=1
CCG007968.1	1476	21	444	0.61135801	10.51590439	4.104411647	Up	1.76E-10	3.72E-10	AAEL002186-PA OS=Aedes aegypti GN=AAEL002186 PE=4 SV=1
CCG005748.1	601	72	444	5.14778851	25.82608134	2.326804069	Up	0	0	Glyoxylase OS=Aedes aegypti PE=2 SV=1
CCG006951.1	1113	94	444	3.62907013	13.94561984	1.942140218	Up	0	0	AAEL003087-PA OS=Aedes aegypti GN=AAEL003087 PE=4 SV=1
CCG019628.1	1882	146	444	3.33346507	8.247329908	1.306904511	Up	9.33E-15	3.33E-14	AAEL004910-PA OS=Aedes aegypti GN=AAEL004910 PE=4 SV=1
CCG019383.1	681	148	444	9.33850326	22.79218045	1.287275705	Up	0	0	AAEL013890-PA OS=Aedes aegypti GN=AAEL013890 PE=4 SV=1
CCG022550.1	670	6	445	0.38480359	23.21855696	5.915012229	Up	8.38E-06	1.48E-05	-

CCG005710.1	1632	20	445	0.52658988	9.532128164	4.178046635	Up	3.69E-10	7.72E-10	AAEL002838-PA OS=Aedes aegypti GN=AAEL002838 PE=4 SV=1
CCG002744.1	299	32	445	4.59876758	52.02820456	3.49997473	Up	1.77E-13	5.17E-13	AAEL001029-PA OS=Aedes aegypti GN=AAEL001029 PE=4 SV=1
CCG003099.1	1215	66	445	2.33415842	12.80364869	2.45558061	Up	0	0	AAEL002170-PA OS=Aedes aegypti GN=AAEL002170 PE=4 SV=1
CCG016399.1	1164	145	445	5.35275904	13.3646333	1.32006564	Up	0	0	AAEL007668-PA OS=Aedes aegypti GN=AAEL007668 PE=4 SV=1
CCG018917.1	2367	149	445	2.70489669	6.572215109	1.280806209	Up	2.33E-13	6.62E-13	AAEL000922-PA OS=Aedes aegypti GN=AAEL000922 PE=4 SV=1
CCG004116.1	1602	174	445	4.66712473	9.710632437	1.057031234	Up	1.78E-13	5.20E-13	AAEL008643-PA OS=Aedes aegypti GN=AAEL008643 PE=4 SV=1
CCG008231.1	2244	1545	445	29.5847772	6.932456847	-2.093416393	Down	7.67E-196	4.57E-195	AAEL003154-PA OS=Aedes aegypti GN=AAEL003154 PE=4 SV=1
CCG028600.1	873	49	446	2.41181786	17.85955491	2.88850326	Up	0	0	Ring finger protein 121 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006831 PE=4 SV=1
CCG020582.1	539	120	446	9.56654574	28.92651473	1.596322508	Up	0	0	AAEL000561-PA OS=Aedes aegypti GN=AAEL000561 PE=4 SV=1
CCG010805.1	2414	129	446	2.29622857	6.458737134	1.491985848	Up	0	0	AAEL013552-PA OS=Aedes aegypti GN=AAEL013552 PE=4 SV=1
CCG027314.1	1929	862	446	19.2016129	8.082629052	-1.248330955	Down	6.98E-53	3.30E-52	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008752 PE=4 SV=1
CCG014539.1	4616	3050	446	28.3920473	3.377684454	-3.071380424	Down	0	0	AAEL009795-PA OS=Aedes aegypti GN=AAEL009795 PE=4 SV=1
CCG004222.1	489	20	447	1.75745336	31.95572539	4.18451613	Up	3.69E-10	7.72E-10	AAEL005232-PA OS=Aedes aegypti GN=AAEL005232 PE=4 SV=1
CCG004899.1	1278	30	447	1.00867922	12.2271907	3.599553629	Up	3.71E-13	9.95E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001997 PE=4 SV=1
CCG017874.1	2004	55	447	1.17930908	7.797579699	2.725084511	Up	0	0	AAEL010355-PA OS=Aedes aegypti GN=AAEL010355 PE=4 SV=1

CCG012166.1	1033	110	447	4.57567358	15.12715365	1.725084511	Up	0	0	Peptidyl-prolyl cis-trans isomerase OS=Aedes aegypti GN=AAEL003477 PE=3 SV=1
CCG015934.1	1223	118	447	4.14589426	12.77706436	1.623801176	Up	0	0	AAEL003084-PA (Fragment) OS=Aedes aegypti GN=AAEL801117_b PE=2 SV=1
CCG017246.1	2304	172	447	3.20781005	6.782269843	1.08017947	Up	1.35E-13	4.09E-13	AAEL010800-PA OS=Aedes aegypti GN=AAEL010800 PE=4 SV=1
CCG003662.1	2547	886	447	14.9474617	6.135198161	-1.284718664	Down	9.75E-57	4.66E-56	AAEL006390-PA OS=Aedes aegypti GN=AAEL006390 PE=2 SV=1
CCG002345.1	1914	1281	447	28.7587409	8.164237052	-1.816610535	Down	1.12E-134	6.27E-134	Putative carboxylesterase OS=Aedes aegypti PE=2 SV=1
CCG028623.1	1449	1427	447	42.3173301	10.78423031	-1.972325394	Down	1.95E-167	1.13E-166	AAEL011981-PA OS=Aedes aegypti GN=AAEL011981 PE=3 SV=1
CCG002946.1	1032	30	448	1.24912019	15.17568604	3.60277753	Up	3.71E-13	9.96E-13	Cytosine-specific methyltransferase OS=Aedes aegypti GN=AAEL006166 PE=3 SV=1
CCG011211.1	786	0	449	0.001	19.96980442	14.28553258	Up	4.44E-16	1.65E-15	AAEL013288-PA OS=Aedes aegypti GN=AAEL013288 PE=4 SV=1
CCG018502.1	990	0	449	0.001	15.85481441	13.95263337	Up	4.44E-16	1.63E-15	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG002828.1	2139	6	449	0.12053221	7.338132899	5.927922338	Up	8.38E-06	1.47E-05	AAEL002320-PA OS=Aedes aegypti GN=AAEL002320 PE=4 SV=1
CCG007399.1	1437	35	449	1.04658365	10.92294104	3.383601822	Up	6.39E-14	2.11E-13	Wd repeat protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009423 PE=4 SV=1
CCG015899.1	1642	121	449	3.16646644	9.559236462	1.594021601	Up	0	0	AAEL006694-PA OS=Aedes aegypti GN=AAEL006694 PE=4 SV=1
CCG007924.1	1275	132	449	4.44863135	12.31079708	1.468490719	Up	1.46E-13	4.36E-13	AAEL006870-PA OS=Aedes aegypti GN=AAEL006870 PE=4 SV=1
CCG013860.1	1722	139	449	3.46852097	9.115137207	1.393943766	Up	1.46E-13	4.37E-13	-
CCG023090.1	707	1082	449	65.7613194	22.20122528	-1.566599945	Down	1.46E-92	7.62E-92	AAEL001325-PA OS=Aedes aegypti GN=AAEL001325 PE=4 SV=1
CCG013241.1	1266	19	450	0.64488543	12.42592776	4.268166882	Up	7.73E-10	1.60E-09	AAEL012371-PA OS=Aedes aegypti GN=AAEL012371 PE=4 SV=1

CCG020735.1	2087	32	450	0.65885554	7.537721393	3.516094395	Up	1.77E-13	5.16E-13	AAEL008101-PA OS=Aedes aegypti GN=AAEL008101 PE=4 SV=1
CCG002710.1	1188	68	450	2.4595471	13.2417715	2.428631554	Up	0	0	AAEL007581-PA OS=Aedes aegypti GN=AAEL007581 PE=4 SV=1
CCG000414.1	1768	73	450	1.77420284	8.897751441	2.326269836	Up	0	0	AAEL003913-PA OS=Aedes aegypti GN=AAEL003913 PE=4 SV=1
CCG010180.1	1800	82	450	1.95751013	8.739569193	2.15854239	Up	0	0	AAEL006453-PA OS=Aedes aegypti GN=AAEL006453 PE=4 SV=1
CCG013109.1	1539	119	450	3.32254608	10.22171835	1.621276632	Up	0	0	AAEL012306-PA OS=Aedes aegypti GN=AAEL012306 PE=4 SV=1
CCG013347.1	1347	124	450	3.95564001	11.67871162	1.561898085	Up	1.61E-13	4.77E-13	AAEL008419-PA OS=Aedes aegypti GN=AAEL008419 PE=4 SV=1
CCG018377.1	1374	130	450	4.06554985	11.44921728	1.493726582	Up	0	0	AAEL001533-PA OS=Aedes aegypti GN=AAEL001533 PE=3 SV=1
CCG011446.1	1458	148	450	4.3618112	10.7895916	1.306641029	Up	0	0	AAEL014348-PB OS=Aedes aegypti GN=CASPS8 PE=4 SV=1
CCG008101.1	930	1225	450	56.5999192	16.91529521	-1.742471639	Down	1.21E-121	6.62E-121	AAEL000757-PB OS=Aedes aegypti GN=AAEL000757 PE=4 SV=1
CCG014618.1	642	13258	450	887.371871	24.50346503	-5.178481142	Down	0	0	AAEL004212-PA OS=Aedes aegypti GN=AAEL004212 PE=4 SV=1
CCG024435.1	2661	35	451	0.56517877	5.924908991	3.39001381	Up	6.39E-14	2.11E-13	AAEL013016-PA OS=Aedes aegypti GN=AAEL013016 PE=4 SV=1
CCG010762.1	717	64	451	3.83551327	21.98909738	2.519296827	Up	0	0	AAEL005132-PA OS=Aedes aegypti GN=AAEL005132 PE=4 SV=1
CCG011128.1	1806	116	451	2.75996081	8.729890822	1.661315832	Up	1.16E-13	3.59E-13	Mitochondrial GTPase 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017444 PE=3 SV=1
CCG020694.1	312	141	451	19.4190147	50.53263726	1.379745475	Up	0	0	-
CCG005579.1	3306	991	451	12.8805224	4.768960322	-1.43344442	Down	1.41E-74	7.05E-74	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006609 PE=4 SV=1
CCG019361.1	2114	1142	451	23.2126002	7.457986199	-1.638050108	Down	4.73E-104	2.52E-103	-
CCG005412.1	610	3455	451	243.377759	25.84620135	-3.235173168	Down	0	0	AAEL002797-PA OS=Aedes aegypti GN=AAEL002797 PE=4 SV=1

CCG000767.1	2088	20	452	0.4115875	7.567596313	4.200564071	Up	3.69E-10	7.71E-10	AAEL001681-PA OS=Aedes aegypti GN=AAEL001681 PE=4 SV=1
CCG001928.1	1692	68	452	1.72691605	9.338735875	2.435029325	Up	0	0	AAEL004456-PA OS=Aedes aegypti GN=AAEL004456 PE=4 SV=1
CCG018691.1	900	142	452	6.77966924	17.55682345	1.372745047	Up	2.62E-13	7.35E-13	-
CCG022690.1	9075	1296	452	6.13650425	1.741172573	-1.817357837	Down	2.63E-136	1.47E-135	AGAP003681-PA OS=Anopheles gambiae GN=AgaP_AGAP003681 PE=4 SV=2
CCG026843.1	1344	6211	452	198.575165	11.75680141	-4.078117686	Down	0	0	AAEL003674-PA OS=Aedes aegypti GN=AAEL003674 PE=4 SV=1
CCG027133.1	1665	17	453	0.43873002	9.511170798	4.438217603	Up	3.39E-09	6.88E-09	AAEL000404-PA OS=Aedes aegypti GN=AAEL000404 PE=4 SV=1
CCG002307.1	829	26	453	1.34766357	19.10265305	3.825240726	Up	4.60E-12	1.05E-11	Ribulose-phosphate 3-epimerase OS=Aedes aegypti PE=2 SV=1
CCG002208.2	1038	36	453	1.49027981	15.25635778	3.355755442	Up	6.13E-14	2.03E-13	AAEL008443-PA OS=Aedes aegypti GN=AAEL008443 PE=4 SV=1
CCG008717.1	942	80	453	3.64923436	16.81114584	2.203752349	Up	0	0	AAEL006646-PA OS=Aedes aegypti GN=AAEL006646 PE=4 SV=1
CCG007850.1	1218	137	453	4.83321317	13.00172363	1.427648361	Up	2.90E-13	8.04E-13	AAEL014441-PA OS=Aedes aegypti GN=AAEL014441 PE=4 SV=1
CCG002730.1	465	1917	453	177.146196	34.05612769	-2.378954178	Down	1.27E-284	7.96E-284	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015665 PE=4 SV=1
CCG021004.1	2586	2335	453	38.799045	6.123781662	-2.663526391	Down	0	0	AAEL014344-PA OS=Aedes aegypti GN=AAEL014344 PE=4 SV=1
CCG009874.1	1614	15	454	0.39934698	9.833369055	4.621971096	Up	1.48E-08	2.93E-08	AAEL007875-PA OS=Aedes aegypti GN=AAEL007875 PE=4 SV=1
CCG002020.1	1414	71	454	2.15760336	11.22422748	2.379114572	Up	0	0	AAEL010811-PA OS=Aedes aegypti GN=AAEL010811 PE=4 SV=1
CCG017619.1	955	107	454	4.81441006	16.61890854	1.787394705	Up	0	0	AAEL014899-PA OS=Aedes aegypti GN=AAEL014899 PE=3 SV=1
CCG010624.1	1696	115	454	2.91363177	9.357934938	1.68337164	Up	6.13E-14	2.03E-13	AAEL001971-PA OS=Aedes aegypti GN=AAEL001971 PE=4 SV=1
CCG012199.1	12315	924	454	3.22403855	1.288758234	-1.32288735	Down	8.90E-62	4.32E-61	AAEL005903-PA OS=Aedes aegypti GN=AAEL005903 PE=4 SV=1

CCG000368.1	930	26	455	1.20130441	17.10324294	3.831596221	Up	4.60E-12	1.05E-11	AAEL003654-PA OS=Aedes aegypti GN=AAEL003654 PE=4 SV=1
CCG000487.1	1599	27	455	0.72556775	9.94747713	3.777148437	Up	2.28E-12	5.42E-12	AAEL003172-PA OS=Aedes aegypti GN=AAEL003172 PE=4 SV=1
CCG022239.1	954	38	455	1.71158272	16.67297267	3.284108425	Up	4.26E-14	1.45E-13	AAEL004455-PA OS=Aedes aegypti GN=AAEL004455 PE=4 SV=1
CCG002890.1	1923	872	455	19.4849759	8.271459143	-1.236148386	Down	1.12E-52	5.27E-52	AAEL011814-PA OS=Aedes aegypti GN=AAEL011814 PE=4 SV=1
CCG022846.1	4911	1682	455	14.716981	3.238854802	-2.183926051	Down	6.87E-225	4.17E-224	AAEL014592-PA OS=Aedes aegypti GN=AAEL014592 PE=4 SV=1
CCG024922.1	2676	22	456	0.35326389	5.957015773	4.075771599	Up	8.38E-11	1.80E-10	AAEL006140-PA OS=Aedes aegypti GN=AAEL006140 PE=3 SV=1
CCG025394.1	2361	60	456	1.09198817	6.751789161	2.628312622	Up	0	0	Ubiquitin conjugating enzyme 7 interacting protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017240 PE=4 SV=1
CCG013792.1	1323	85	456	2.76071613	12.04911127	2.125812282	Up	0	0	AAEL013391-PA OS=Aedes aegypti GN=AAEL013391 PE=4 SV=1
CCG026959.1	1214	90	456	3.18556517	13.13095075	2.043350122	Up	0	0	AAEL005070-PA OS=Aedes aegypti GN=AAEL005070 PE=4 SV=1
CCG004794.1	1614	126	456	3.3545146	9.876687861	1.557923294	Up	1.34E-13	4.06E-13	AAEL011017-PA OS=Aedes aegypti GN=AAEL011017 PE=4 SV=1
CCG012847.1	1194	127	456	4.57048266	13.35089967	1.546518531	Up	1.57E-13	4.67E-13	AAEL011058-PA (Fragment) OS=Aedes aegypti GN=AAEL011058 PE=4 SV=1
CCG026580.1	2244	169	456	3.2361342	7.103820948	1.134323782	Up	0	0	AAEL003147-PA OS=Aedes aegypti GN=AAEL003147 PE=4 SV=1
CCG012658.1	1164	48	457	1.77194782	13.72502791	2.953401058	Up	5.35E-14	1.79E-13	AAEL001917-PA OS=Aedes aegypti GN=AAEL001917 PE=4 SV=1
CCG013914.1	1337	99	457	3.18175297	11.94908937	1.909006939	Up	0	0	AAEL002808-PA OS=Aedes aegypti GN=AAEL002808 PE=3 SV=1
CCG020697.1	1407	113	457	3.45101635	11.35460731	1.718184596	Up	1.42E-13	4.26E-13	AAEL009727-PA OS=Aedes aegypti GN=AAEL009727 PE=4 SV=1
CCG026602.1	1471	145	457	4.23562986	10.86059312	1.358454469	Up	0	0	AAEL003822-PA OS=Aedes aegypti GN=AAEL003822 PE=4 SV=1

CCG006553.1	1872	155	457	3.5578573	8.534151968	1.262239154	Up	3.42E-13	9.27E-13	AAEL010006-PA OS=Aedes aegypti GN=AAEL010006 PE=4 SV=1
CCG017773.1	786	109	458	5.95890722	20.37009003	1.773332667	Up	0	0	Activin receptor type I OS=Culex quinquefasciatus GN=CpipJ_CPIJ017088 PE=4 SV=1
CCG020079.1	3109	151	458	2.08698293	5.149852287	1.303112253	Up	0	0	AAEL011162-PA OS=Aedes aegypti GN=AAEL011162 PE=4 SV=1
CCG000948.1	603	2399	458	170.952559	26.55205765	-2.68670045	Down	0	0	AAEL008166-PA OS=Aedes aegypti GN=AAEL008166 PE=3 SV=1
CCG019598.1	702	60	459	3.67262689	22.85733481	2.637772952	Up	0	0	AAEL007351-PA OS=Aedes aegypti GN=AAEL007351 PE=4 SV=1
CCG011422.1	1824	84	459	1.97886936	8.797066359	2.152346124	Up	0	0	AAEL005297-PA OS=Aedes aegypti GN=AAEL005297 PE=4 SV=1
CCG018211.1	972	122	459	5.3933206	16.50807514	1.61392621	Up	8.04E-14	2.58E-13	AAEL002939-PA OS=Aedes aegypti GN=AAEL002939 PE=4 SV=1
CCG025694.1	357	184	459	22.146866	44.94635585	1.021101591	Up	3.33E-13	9.08E-13	AAEL003764-PA OS=Aedes aegypti GN=AAEL003764 PE=4 SV=1
CCG011617.1	270	843	459	134.16106	59.42907051	-1.174725274	Down	3.86E-47	1.78E-46	Activating transcription factor OS=Aedes aegypti PE=2 SV=1
CCG017906.1	2157	34	460	0.67731617	7.455172608	3.460340414	Up	8.82E-14	2.79E-13	DNA mismatch repair protein spellchecker 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017719 PE=3 SV=1
CCG027796.1	1194	58	460	2.08730704	13.46801283	2.68982226	Up	0	0	AAEL001321-PA OS=Aedes aegypti GN=AAEL001321 PE=3 SV=1
CCG017543.1	564	65	460	4.95218573	28.51206971	2.525435442	Up	0	0	D-tyrosyl-tRNA(Tyr) deacylase OS=Aedes aegypti GN=AAEL014489 PE=3 SV=1
CCG014307.1	2698	122	460	1.9430347	5.960269576	1.617065917	Up	8.04E-14	2.58E-13	AAEL010559-PA OS=Aedes aegypti GN=AAEL010559 PE=4 SV=1
CCG014662.1	477	180	460	16.2149942	33.71238431	1.055950158	Up	3.32E-13	9.05E-13	AAEL014545-PA OS=Aedes aegypti GN=AAEL014545 PE=4 SV=1
CCG027345.2	549	768	460	60.110667	29.29108801	-1.037159246	Down	1.69E-35	7.33E-35	AAEL000288-PA OS=Aedes aegypti GN=AAEL000288 PE=4 SV=1

CCG009803.1	33896	1059	460	1.34248728	0.474416076	-1.500683619	Down	3.89E-85	2.00E-84	AAEL005977-PA OS=Aedes aegypti GN=AAEL005977 PE=4 SV=1
CCG020538.1	752	1067	460	60.969025	21.38405228	-1.511541206	Down	1.13E-86	5.80E-86	AAEL012352-PA OS=Aedes aegypti GN=AAEL012352 PE=4 SV=1
CCG028211.1	3294	2824	460	36.8386553	4.881848001	-2.915721119	Down	0	0	-
CCG024623.1	489	3	461	0.263618	32.95657585	6.965973644	Up	5.28E-05	8.84E-05	Predicted protein OS=Nematostella vectensis GN=v1g229551 PE=3 SV=1
CCG024252.1	804	8	461	0.42755955	20.04448457	5.550936144	Up	2.19E-06	3.97E-06	AAEL008084-PA OS=Aedes aegypti GN=AAEL008084 PE=4 SV=1
CCG015694.1	1413	13	461	0.39533372	11.40535428	4.850496426	Up	6.34E-08	1.23E-07	DNA primase OS=Aedes aegypti GN=AAEL014314 PE=3 SV=1
CCG003819.1	807	17	461	0.90518648	19.96996975	4.463473303	Up	3.39E-09	6.88E-09	AAEL004960-PA OS=Aedes aegypti GN=AAEL004960 PE=4 SV=1
CCG005578.1	1953	33	461	0.7260631	8.251800098	3.506542025	Up	1.31E-13	4.01E-13	AAEL007882-PA OS=Aedes aegypti GN=AAEL007882 PE=4 SV=1
CCG005961.1	2040	50	461	1.05317977	7.899885094	2.907079954	Up	0	0	AAEL015662-PA OS=Aedes aegypti GN=AAEL015662 PE=4 SV=1
CCG017263.1	873	170	461	8.36753137	18.46021259	1.141545208	Up	3.53E-13	9.52E-13	AAEL006867-PA OS=Aedes aegypti GN=AAEL006867 PE=4 SV=1
CCG009112.1	1127	934	461	35.6111199	14.29970328	-1.316342595	Down	5.85E-62	2.84E-61	-
CCG009181.1	693	32	462	1.98417245	23.30551785	3.554062245	Up	1.77E-13	5.16E-13	AAEL009789-PA OS=Aedes aegypti GN=AAEL009789 PE=4 SV=1
CCG002900.1	1650	56	462	1.45836675	9.788317496	2.746707323	Up	0	0	AAEL012679-PA OS=Aedes aegypti GN=AAEL012679 PE=4 SV=1
CCG016578.1	937	76	462	3.48527196	17.23663166	2.306134732	Up	0	0	AAEL010083-PA OS=Aedes aegypti GN=IMD PE=4 SV=1
CCG027802.1	2494	98	462	1.68846591	6.475831543	1.939352401	Up	0	0	AAEL003170-PA OS=Aedes aegypti GN=AAEL003170 PE=4 SV=1
CCG002477.3	1764	156	462	3.80004456	9.155739155	1.268660026	Up	1.42E-13	4.27E-13	AAEL009764-PB OS=Aedes aegypti GN=AAEL009764 PE=4 SV=1
CCG014367.1	510	940	462	79.1991187	31.66808602	-1.322454701	Down	8.75E-63	4.25E-62	AAEL014863-PA OS=Aedes aegypti GN=AAEL014863 PE=4 SV=1
CCG010544.1	1287	1261	462	42.1016592	12.549125	-1.746290315	Down	1.52E-125	8.37E-125	AAEL014318-PA OS=Aedes aegypti GN=AAEL014318 PE=4 SV=1

CCG002490.1	477	7	463	0.63058311	33.93224768	5.749826665	Up	4.31E-06	7.69E-06	-
CCG021695.1	1848	23	463	0.53479648	8.75848601	4.033619631	Up	4.01E-11	8.66E-11	AAEL000938-PA OS=Aedes aegypti GN=AAEL000938 PE=4 SV=1
CCG008117.1	495	46	463	3.99314705	32.69834777	3.033619631	Up	3.40E-14	1.18E-13	AAEL011767-PA OS=Aedes aegypti GN=AAEL011767 PE=4 SV=1
CCG012559.1	1035	53	463	2.20038255	15.63834024	2.829261133	Up	0	0	AAEL011684-PA OS=Aedes aegypti GN=AAEL011684 PE=4 SV=1
CCG008181.1	780	137	463	7.54724826	20.75087455	1.459149504	Up	2.90E-13	8.04E-13	AAEL006290-PA OS=Aedes aegypti GN=AAEL006290 PE=4 SV=1
CCG012339.1	1587	140	463	3.79065082	10.19891755	1.42789857	Up	0	0	AAEL007810-PA OS=Aedes aegypti GN=CYP4D23 PE=3 SV=2
CCG026111.1	2601	147	463	2.42850865	6.222868953	1.357509242	Up	7.02E-14	2.29E-13	AAEL013943-PB (Fragment) OS=Aedes aegypti GN=AAEL013943 PE=4 SV=1
CCG007251.1	23177	861	463	1.59627827	0.698351044	-1.19268784	Down	2.99E-49	1.39E-48	AAEL010879-PA OS=Aedes aegypti GN=AAEL010879 PE=4 SV=1
CCG020015.2	1452	1324	463	39.181769	11.14716401	-1.81350582	Down	8.36E-139	4.70E-138	AAEL008309-PA OS=Aedes aegypti GN=AAEL008309 PE=4 SV=1
CCG011763.1	981	1573	463	68.9005021	16.49916631	-2.062121368	Down	1.70E-195	1.01E-194	AAEL006966-PA OS=Aedes aegypti GN=AAEL006966 PE=3 SV=1
CCG027449.1	1979	1641	463	35.6307905	8.178717608	-2.123177936	Down	1.06E-211	6.34E-211	AAEL000119-PA OS=Aedes aegypti GN=AAEL000119 PE=4 SV=1
CCG019874.1	1914	3648	463	81.8984283	8.456469251	-3.275708427	Down	0	0	AAEL003434-PA OS=Aedes aegypti GN=AAEL003434 PE=4 SV=1
CCG002444.1	2242	46	464	0.88162703	7.234897601	3.036732243	Up	3.40E-14	1.18E-13	AAEL011047-PA (Fragment) OS=Aedes aegypti GN=AAEL011047 PE=4 SV=1
CCG005012.1	1551	110	464	3.04749891	10.45818209	1.778934485	Up	0	0	AAEL008547-PA OS=Aedes aegypti GN=AAEL008547 PE=4 SV=1
CCG016105.1	804	121	464	6.46683817	20.1749259	1.641430962	Up	0	0	Chip-domain serine protease OS=Culex quinquefasciatus GN=CpipJ_CPIJ000615 PE=3 SV=1
CCG005444.1	1317	138	464	4.50252344	12.31635567	1.451769742	Up	1.49E-13	4.45E-13	AAEL003885-PA OS=Aedes aegypti GN=AAEL003885 PE=4 SV=1

CCG005124.1	485	153	464	13.5554008	33.44461943	1.302906356	Up	0	0	AAEL003985-PA OS=Aedes aegypti GN=AAEL003985 PE=4 SV=1
CCG008832.1	648	174	464	11.5381695	25.0318525	1.117350703	Up	1.78E-13	5.19E-13	AAEL004425-PA OS=Aedes aegypti GN=AAEL004425 PE=4 SV=1
CCG005021.1	1031	177	464	7.37695735	15.73291991	1.092688649	Up	0	0	AAEL013113-PA OS=Aedes aegypti GN=AAEL013113 PE=4 SV=1
CCG001045.1	705	62	465	3.77889865	23.05758681	2.609203799	Up	0	0	AAEL006646-PA OS=Aedes aegypti GN=AAEL006646 PE=4 SV=1
CCG018974.1	1569	68	465	1.8622957	10.36048356	2.475937269	Up	0	0	AAEL012628-PA OS=Aedes aegypti GN=AAEL012628 PE=4 SV=1
CCG000367.1	1535	94	465	2.63137137	10.58996658	2.008811258	Up	0	0	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1
CCG018719.1	1224	137	465	4.80952095	13.28071789	1.465368027	Up	2.90E-13	8.04E-13	AAEL005368-PA OS=Aedes aegypti GN=AAEL005368 PE=4 SV=1
CCG022181.1	1664	1592	465	41.1104672	9.768989603	-2.073224511	Down	3.21E-199	1.92E-198	AAEL007218-PA OS=Aedes aegypti GN=AAEL007218 PE=4 SV=1
CCG023975.1	1203	8	466	0.28575052	13.54161012	5.566499348	Up	2.19E-06	3.96E-06	AAEL007000-PA OS=Aedes aegypti GN=AAEL007000 PE=4 SV=1
CCG024058.1	2240	42	466	0.80568252	7.272570079	3.174181926	Up	7.66E-14	2.47E-13	AAEL013014-PA OS=Aedes aegypti GN=AAEL013014 PE=4 SV=1
CCG017816.1	702	158	466	9.67125081	23.20592162	1.2627186	Up	0	0	AAEL005915-PA OS=Aedes aegypti GN=AAEL005915 PE=4 SV=1
CCG023779.1	1648	955	466	24.900544	9.885046709	-1.332857574	Down	1.62E-64	7.90E-64	AAEL001645-PA OS=Aedes aegypti GN=AAEL001645 PE=4 SV=1
CCG005848.1	10089	2447	466	10.4219388	1.614685001	-2.690299038	Down	0	0	AAEL001184-PA OS=Aedes aegypti GN=AAEL001184 PE=4 SV=1
CCG017831.1	192	6	467	1.34280421	85.02872527	5.984629443	Up	8.38E-06	1.47E-05	Putative salivary basic peptide 4.2k-1 OS=Aedes albopictus PE=4 SV=1
CCG013394.1	792	33	467	1.79040561	20.61302431	3.525197824	Up	1.31E-13	4.01E-13	Glyoxylase 3 OS=Aedes aegypti PE=2 SV=1
CCG020190.1	1155	94	467	3.49710394	14.13464524	2.015003092	Up	0	0	AAEL012784-PA OS=Aedes aegypti GN=AAEL012784 PE=4 SV=1
CCG012855.1	1314	126	467	4.12038551	12.42428862	1.59231202	Up	1.34E-13	4.06E-13	AAEL003415-PA OS=Aedes aegypti GN=AAEL003415 PE=4 SV=1

CCG022462.1	2676	55	468	0.88315972	6.113779346	2.79131821	Up	0	0	AAEL004136-PA OS=Aedes aegypti GN=AAEL004136 PE=4 SV=1
CCG012896.1	1235	75	468	2.60949805	13.24734699	2.343859233	Up	0	0	AAEL009502-PA OS=Aedes aegypti GN=AAEL009502 PE=4 SV=1
CCG007547.1	723	130	468	7.72623167	22.62859409	1.55031011	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_22817 PE=4 SV=1
CCG010300.1	2190	132	468	2.58995661	7.470535858	1.528283804	Up	1.46E-13	4.36E-13	AAEL012505-PA OS=Aedes aegypti GN=AAEL012505 PE=4 SV=1
CCG011959.1	2319	189	468	3.50206116	7.05496918	1.010435499	Up	2.34E-13	6.65E-13	AAEL012975-PA OS=Aedes aegypti GN=AAEL012975 PE=4 SV=1
CCG022104.1	449	1017	468	97.3278844	36.43758024	-1.41742604	Down	3.21E-75	1.61E-74	AAEL007764-PA OS=Aedes aegypti GN=AAEL007764 PE=4 SV=1
CCG009881.1	938	1238	468	56.7127201	17.44186943	-1.701117676	Down	7.68E-119	4.20E-118	AAEL010955-PA OS=Aedes aegypti GN=AAEL010955 PE=4 SV=1
CCG009401.1	1557	1252	468	34.5524134	10.50769013	-1.717340924	Down	8.86E-122	4.86E-121	AAEL011677-PA OS=Aedes aegypti GN=AAEL011677 PE=4 SV=1
CCG017485.1	540	8	469	0.63658866	30.36191075	5.575757316	Up	2.19E-06	3.96E-06	Putative salivary protein 34k-1 OS=Aedes albopictus PE=2 SV=1
CCG026337.1	1165	35	469	1.29093623	14.07333202	3.446474299	Up	6.39E-14	2.11E-13	-
CCG027985.1	1512	50	469	1.42095683	10.84353955	2.931901127	Up	0	0	AAEL001008-PA OS=Aedes aegypti GN=AAEL001008 PE=4 SV=1
CCG014679.1	2559	60	469	1.00749671	6.406968271	2.668866721	Up	0	0	AAEL000707-PA OS=Aedes aegypti GN=AAEL000707 PE=4 SV=1
CCG019160.1	867	148	469	7.33508734	18.91053265	1.366303951	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011649 PE=4 SV=1
CCG005531.1	1035	182	469	7.55603063	15.84099691	1.067962676	Up	0	0	AAEL006868-PA OS=Aedes aegypti GN=AAEL006868 PE=3 SV=1
CCG008642.1	1952	876	469	19.2835489	8.399299081	-1.199029743	Down	1.73E-50	8.06E-50	-
CCG016415.1	1683	2036	469	51.9824003	9.741789546	-2.41576453	Down	8.00E-308	5.06E-307	Purine nucleoside phosphorylase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007359 PE=3 SV=1
CCG006332.1	2484	38	470	0.65734699	6.614488761	3.330902637	Up	4.26E-14	1.45E-13	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1

CCG023704.1	537	57	470	4.56103328	30.59662958	2.745940136	Up	0	0	AAEL007838-PA OS=Aedes aegypti GN=AAEL007838 PE=4 SV=1
CCG005109.1	2150	111	470	2.21843746	7.642041899	1.784414284	Up	0	0	AAEL006013-PA OS=Aedes aegypti GN=AAEL006013 PE=4 SV=1
CCG018243.1	355	118	470	14.2828977	46.28278897	1.696187101	Up	0	0	AAEL013457-PA OS=Aedes aegypti GN=AAEL013457 PE=4 SV=1
CCG002103.1	3120	147	470	2.02453557	5.266150668	1.379157806	Up	7.02E-14	2.29E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003554 PE=4 SV=1
CCG013975.1	747	13	471	0.74779993	22.04196568	4.881456735	Up	6.34E-08	1.23E-07	AAEL010421-PA OS=Aedes aegypti GN=AAEL010421 PE=4 SV=1
CCG006786.1	1431	40	471	1.20111068	11.50618334	3.259968359	Up	6.86E-14	2.25E-13	Ryndremine 5-monooxygenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ017147 PE=4 SV=1
CCG017659.2	2324	66	471	1.22031088	7.084917539	2.537502334	Up	0	0	AAEL000853-PA OS=Aedes aegypti GN=AAEL000853 PE=4 SV=1
CCG005084.1	678	100	471	6.33771897	24.28517457	1.938040264	Up	0	0	AAEL005512-PA OS=Aedes aegypti GN=AAEL005512 PE=4 SV=1
CCG007306.1	3723	112	471	1.29266996	4.422602299	1.774541531	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008434 PE=4 SV=1
CCG004826.1	966	116	471	5.15992672	17.04487408	1.723915458	Up	1.16E-13	3.59E-13	AAEL010099-PA OS=Aedes aegypti GN=AAEL010099 PE=4 SV=1
CCG019629.2	621	773	471	53.4872864	26.51424857	-1.01242815	Down	2.19E-34	9.44E-34	AAEL004902-PA OS=Aedes aegypti GN=AAEL004902 PE=3 SV=1
CCG020629.1	4392	54	472	0.52831641	3.756900418	2.830068751	Up	0	0	AAEL005539-PA OS=Aedes aegypti GN=AAEL005539 PE=4 SV=1
CCG015764.1	1503	72	472	2.0584304	10.97824793	2.415031252	Up	0	0	AAEL014410-PA OS=Aedes aegypti GN=AAEL014410 PE=4 SV=1
CCG009303.1	1569	106	472	2.90299036	10.51644782	1.857035799	Up	0	0	AAEL007778-PA OS=Aedes aegypti GN=AAEL007778 PE=4 SV=1
CCG011836.1	1767	108	472	2.62633352	9.338034316	1.830068751	Up	0	0	AAEL004866-PA OS=Aedes aegypti GN=AAEL004866 PE=4 SV=1
CCG001537.1	2104	179	472	3.6556951	7.842351063	1.101140476	Up	0	0	AAEL009500-PA OS=Aedes aegypti GN=AAEL009500 PE=4 SV=1

CCG019931.1	3507	1556	472	19.0649863	4.704963398	-2.018670092	Down	4.37E-188	2.58E-187	AAEL009821-PA OS=Aedes aegypti GN=AAEL009821 PE=4 SV=1
CCG006331.1	1470	40	473	1.16924448	11.24847953	3.266081482	Up	6.86E-14	2.24E-13	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1
CCG028222.1	861	48	473	2.39552527	19.20472115	3.003047076	Up	5.35E-14	1.79E-13	AAEL010253-PA OS=Aedes aegypti GN=AAEL010253 PE=4 SV=1
CCG016040.1	1020	71	473	2.99103055	16.21104403	2.438262458	Up	0	0	AAEL003455-PA OS=Aedes aegypti GN=AAEL003455 PE=4 SV=1
CCG028422.2	3822	133	473	1.49528381	4.326338282	1.532727142	Up	9.90E-14	3.11E-13	AAEL001410-PA (Fragment) OS=Aedes aegypti GN=AAEL001410 PE=4 SV=1
CCG010863.1	729	2	474	0.11788679	22.73007296	7.591056453	Up	9.36E-05	0.00015387	NOS protein OS=Aedes aegypti GN=nos PE=2 SV=1
CCG026299.1	1228	104	474	3.63913062	13.49366709	1.890616735	Up	0	0	AAEL013314-PA OS=Aedes aegypti GN=AAEL013314 PE=4 SV=1
CCG003275.1	1200	117	474	4.18954912	13.80851933	1.720691733	Up	9.44E-14	2.97E-13	AAEL007896-PA (Fragment) OS=Aedes aegypti GN=AAEL007896 PE=4 SV=1
CCG027365.1	3556	142	474	1.71588929	4.659792798	1.441309333	Up	2.62E-13	7.35E-13	AAEL005272-PA OS=Aedes aegypti GN=AAEL005272 PE=4 SV=1
CCG009515.2	1206	174	474	6.19961345	13.73982022	1.148112957	Up	1.78E-13	5.20E-13	AAEL001378-PA OS=Aedes aegypti GN=AAEL001378 PE=4 SV=1
CCG007421.1	2265	1523	474	28.8931151	7.315771828	-1.981643774	Down	1.09E-179	6.37E-179	AAEL004086-PA OS=Aedes aegypti GN=AAEL004086 PE=4 SV=1
CCG011318.1	627	15	475	1.02798408	26.48354301	4.687206311	Up	1.48E-08	2.92E-08	Putative 23.4 kDa salivary protein OS=Aedes albopictus PE=2 SV=1
CCG022997.1	1446	44	475	1.30751613	11.48352799	3.134665288	Up	3.91E-14	1.35E-13	AAEL005068-PA OS=Aedes aegypti GN=AAEL005068 PE=4 SV=1
CCG026331.1	2175	89	475	1.75830178	7.634566192	2.118363476	Up	0	0	AAEL009731-PA OS=Aedes aegypti GN=AAEL009731 PE=4 SV=1
CCG003715.1	2061	178	475	3.7111173	8.056856607	1.118363476	Up	1.15E-13	3.57E-13	AAEL002151-PA OS=Aedes aegypti GN=AAEL002151 PE=4 SV=1
CCG004359.1	252	802	475	136.752886	65.89357725	-1.053361519	Down	6.56E-38	2.89E-37	AAEL008785-PA OS=Aedes aegypti GN=AAEL008785 PE=4 SV=1

CCG008816.1	3435	40	476	0.50037537	4.844291046	3.275202872	Up	6.86E-14	2.24E-13	AAEL007237-PA OS=Aedes aegypti GN=AAEL007237 PE=4 SV=1
CCG018429.1	387	88	476	9.77089573	42.99777712	2.137699348	Up	2.93E-14	1.02E-13	AAEL006557-PA OS=Aedes aegypti GN=AAEL006557 PE=4 SV=1
CCG007579.1	741	121	476	7.01665032	22.45632894	1.67826773	Up	0	0	AAEL002459-PA OS=Aedes aegypti GN=AAEL002459 PE=4 SV=1
CCG007769.1	364	21	477	2.47902315	45.81070885	4.207841237	Up	1.76E-10	3.72E-10	-
CCG010803.1	1710	92	477	2.31182198	9.75151931	2.076596704	Up	1.44E-14	5.11E-14	AAEL013548-PA OS=Aedes aegypti GN=AAEL013548 PE=4 SV=1
CCG027354.1	381	98	477	11.0525827	43.76666147	1.985448816	Up	0	0	AAEL005909-PA OS=Aedes aegypti GN=AAEL005909 PE=4 SV=1
CCG018510.1	549	153	477	11.9751719	30.37358474	1.342770817	Up	0	0	AAEL010361-PA OS=Aedes aegypti GN=AAEL010361 PE=4 SV=1
CCG018982.1	1079	154	477	6.13284442	15.45421503	1.333372119	Up	1.82E-13	5.28E-13	AAEL010591-PA OS=Aedes aegypti GN=GSTD6 PE=3 SV=1
CCG016126.1	1314	26	478	0.85023828	12.71693782	3.902740294	Up	4.60E-12	1.05E-11	AAEL006750-PA OS=Aedes aegypti GN=AAEL006750 PE=4 SV=1
CCG020652.1	2574	27	478	0.45073148	6.491863363	3.84829251	Up	2.28E-12	5.41E-12	AAEL010080-PA OS=Aedes aegypti GN=AAEL010080 PE=4 SV=1
CCG025929.1	1854	135	478	3.12886417	9.012975349	1.526364415	Up	1.90E-13	5.49E-13	AAEL000639-PA OS=Aedes aegypti GN=AAEL000639 PE=3 SV=1
CCG005893.1	4131	172	478	1.78910539	4.045039046	1.176915257	Up	1.35E-13	4.09E-13	AAEL012549-PA (Fragment) OS=Aedes aegypti GN=AAEL012549 PE=4 SV=1
CCG001703.1	2149	15	479	0.29992835	7.792003059	4.699304454	Up	1.48E-08	2.93E-08	AAEL009411-PA OS=Aedes aegypti GN=AAEL009411 PE=4 SV=1
CCG018876.1	1030	63	479	2.6282459	16.2572957	2.628915126	Up	0	0	-
CCG028533.1	1335	71	479	2.28528177	12.54308208	2.45644793	Up	0	0	AAEL007318-PA OS=Aedes aegypti GN=AAEL007318 PE=4 SV=1
CCG000154.1	2273	78	479	1.47454435	7.366922382	2.320792831	Up	0	0	AAEL001250-PA OS=Aedes aegypti GN=AAEL001250 PE=4 SV=1
CCG015894.1	921	83	479	3.87240822	18.18134047	2.231155618	Up	1.32E-13	4.02E-13	AAEL006697-PA OS=Aedes aegypti GN=AAEL006697 PE=4 SV=1
CCG015617.1	1065	113	479	4.55923006	15.72301838	1.786016087	Up	1.42E-13	4.27E-13	AAEL007857-PA OS=Aedes aegypti GN=AAEL007857 PE=4 SV=1

CCG007337.1	784	124	479	6.79623354	21.35843696	1.651998739	Up	1.61E-13	4.76E-13	-
CCG003684.1	3213	134	479	1.79207732	5.211644748	1.540105859	Up	0	0	AGAP009234-PA (Fragment) OS=Anopheles gambiae GN=AGAP009234 PE=3 SV=4
CCG005170.1	1449	150	479	4.44821269	11.55625574	1.377376359	Up	0	0	AAEL015402-PA OS=Aedes aegypti GN=AAEL015402 PE=4 SV=1
CCG001073.1	2158	64	480	1.27435728	7.775705677	2.609203799	Up	0	0	AAEL005433-PA OS=Aedes aegypti GN=AAEL005433 PE=4 SV=1
CCG027275.1	996	89	480	3.83966504	16.8473623	2.133470368	Up	0	0	AAEL002282-PA OS=Aedes aegypti GN=AAEL002282 PE=4 SV=1
CCG013618.1	2340	130	480	2.38720748	7.170928569	1.586835986	Up	0	0	AAEL007610-PA OS=Aedes aegypti GN=AAEL007610 PE=4 SV=1
CCG015471.1	2257	142	480	2.70345694	7.434635734	1.45945668	Up	2.62E-13	7.34E-13	AAEL001151-PA OS=Aedes aegypti GN=AAEL001151 PE=3 SV=1
CCG004268.1	1632	155	480	4.08107161	10.28184611	1.333079394	Up	3.42E-13	9.25E-13	AAEL011925-PA OS=Aedes aegypti GN=AAEL011925 PE=4 SV=1
CCG019285.1	1128	156	480	5.94262287	14.87586246	1.323801581	Up	1.42E-13	4.26E-13	AAEL001656-PA OS=Aedes aegypti GN=AAEL001656 PE=4 SV=1
CCG021800.1	6177	1792	480	12.4658838	2.716524664	-2.198151123	Down	1.94E-241	1.19E-240	AAEL000239-PA OS=Aedes aegypti GN=AAEL000239 PE=4 SV=1
CCG027393.1	1896	2270	480	51.4458321	8.850196651	-2.539272783	Down	0	0	AAEL001232-PA OS=Aedes aegypti GN=AAEL001232 PE=3 SV=1
CCG011491.1	2742	3235	480	50.6955111	6.119610814	-3.050346197	Down	0	0	AAEL006085-PA OS=Aedes aegypti GN=AAEL006085 PE=3 SV=1
CCG007081.1	1488	9	481	0.25989759	11.30035694	5.442281286	Up	1.09E-06	2.01E-06	AAEL009709-PA OS=Aedes aegypti GN=AAEL009709 PE=4 SV=1
CCG006232.1	1112	28	481	1.08197173	15.12134094	3.804851366	Up	1.17E-12	2.86E-12	AAEL009918-PA OS=Aedes aegypti GN=AAEL009918 PE=4 SV=1
CCG027389.1	1602	29	481	0.77785412	10.49621169	3.754225292	Up	6.66E-13	1.69E-12	AAEL002849-PA OS=Aedes aegypti GN=AAEL002849 PE=4 SV=1
CCG009394.1	669	82	481	5.26684341	25.1344262	2.254654283	Up	0	0	AAEL008582-PA OS=Aedes aegypti GN=AAEL008582 PE=4 SV=1
CCG000911.1	1641	108	481	2.82798985	10.24675876	1.857318785	Up	0	0	AAEL013048-PA OS=Aedes aegypti GN=AAEL013048 PE=4 SV=1

CCG025611.1	2067	1414	481	29.3948741	8.134944909	-1.853360117	Down	1.37E-152	7.82E-152	AAEL011520-PA OS=Aedes aegypti GN=AAEL011520 PE=4 SV=1
CCG025340.1	1014	53	482	2.2459526	16.61724793	2.887282085	Up	0	0	AAEL012397-PA OS=Aedes aegypti GN=AAEL012397 PE=4 SV=1
CCG013658.1	2169	85	482	1.68392229	7.768505949	2.205811604	Up	0	0	AAEL000125-PA OS=Aedes aegypti GN=AAEL006125 PE=4 SV=1
CCG005815.1	343	814	482	101.974822	49.12504199	-1.053682444	Down	1.81E-38	8.00E-38	AAEL000958-PA OS=Aedes aegypti GN=AAEL000958 PE=4 SV=1
CCG017584.1	2791	1199	482	18.4595886	6.037223004	-1.612413403	Down	1.03E-106	5.52E-106	AAEL002123-PA (Fragment) OS=Aedes aegypti GN=AAEL002123 PE=4 SV=1
CCG013553.1	1338	10	483	0.32114899	12.61946762	5.296264488	Up	5.43E-07	1.01E-06	AAEL000299-PA (Fragment) OS=Aedes aegypti GN=AAEL000299 PE=4 SV=1
CCG025762.1	1984	44	483	0.95295782	8.510507904	3.158760964	Up	3.91E-14	1.35E-13	Zinc finger protein 605 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005857 PE=4 SV=1
CCG028194.1	1482	45	483	1.30474903	11.39328454	3.126339486	Up	0	0	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018272 PE=4 SV=1
CCG016738.2	2160	71	483	1.41243109	7.817059112	2.468445463	Up	0	0	AAEL002404-PA OS=Aedes aegypti GN=AAEL002404 PE=3 SV=1
CCG011392.1	816	93	483	4.89728593	20.6922153	2.079033772	Up	4.04E-14	1.39E-13	AAEL000645-PA OS=Aedes aegypti GN=AAEL000645 PE=4 SV=1
CCG027675.1	845	183	483	9.30587152	19.98206826	1.102492744	Up	3.05E-13	8.42E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017674 PE=4 SV=1
CCG019970.1	954	45	484	2.02687427	17.73564566	3.129323345	Up	0	0	AAEL010791-PA OS=Aedes aegypti GN=AAEL010791 PE=4 SV=1
CCG014396.1	1302	127	484	4.19136428	12.99524267	1.632491754	Up	1.57E-13	4.67E-13	Seryl-tRNA synthetase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007184 PE=3 SV=1
CCG027166.1	2123	145	484	2.93481466	7.96976258	1.441267351	Up	0	0	AAEL015129-PA OS=Aedes aegypti GN=AAEL015129 PE=4 SV=1

CCG017749.1	2090	1433	484	29.4620238	8.095600937	-1.863646453	Down	8.75E-156	5.00E-155	AAEL000378-PA OS=Aedes aegypti GN=AAEL000378 PE=4 SV=1
CCG003677.1	1678	1684	484	43.1233809	10.08331702	-2.096499982	Down	1.10E-213	6.59E-213	Putative nucleoside phosphorylase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007359 PE=3 SV=1
CCG024300.1	1479	132	485	3.83502702	11.4636675	1.579760022	Up	1.46E-13	4.36E-13	AAEL010803-PA (Fragment) OS=Aedes aegypti GN=AAEL010803 PE=3 SV=1
CCG026318.1	701	140	485	8.58168737	24.18653956	1.494871124	Up	0	0	AAEL003749-PA OS=Aedes aegypti GN=AAEL003749 PE=4 SV=1
CCG002117.1	1332	163	485	5.25830836	12.72880198	1.275425987	Up	1.80E-13	5.24E-13	AAEL003885-PA OS=Aedes aegypti GN=AAEL003885 PE=4 SV=1
CCG000714.1	1572	189	485	5.16620855	10.78547343	1.061911717	Up	2.34E-13	6.64E-13	AAEL014582-PA OS=Aedes aegypti GN=AAEL014582 PE=4 SV=1
CCG026562.1	1641	935	485	24.4830602	10.33197089	-1.244668414	Down	5.94E-57	2.84E-56	-
CCG016998.1	837	8	486	0.41070236	20.29835425	5.627125707	Up	2.19E-06	3.96E-06	AAEL012659-PA OS=Aedes aegypti GN=AAEL012659 PE=4 SV=1
CCG001280.1	2835	37	486	0.5608043	5.992847447	3.417672342	Up	5.17E-14	1.74E-13	AAEL003036-PA OS=Aedes aegypti GN=AAEL003036 PE=4 SV=1
CCG011946.1	1508	46	486	1.31074787	11.26639424	3.103563751	Up	3.40E-14	1.18E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003105 PE=4 SV=1
CCG014232.1	1506	119	486	3.39535088	11.28135625	1.732307944	Up	0	0	AAEL000788-PA OS=Aedes aegypti GN=AAEL008788 PE=4 SV=1
CCG017922.1	771	163	486	9.08439266	22.03595656	1.278397553	Up	1.80E-13	5.25E-13	AAEL015052-PA OS=Aedes aegypti GN=AAEL015052 PE=4 SV=1
CCG024900.2	2808	1566	486	23.9638905	6.05047098	-1.98574279	Down	3.25E-185	1.92E-184	AAEL000200-PA OS=Aedes aegypti GN=AAEL000200 PE=4 SV=1
CCG017095.1	942	2544	486	116.045653	18.03579884	-2.685757248	Down	0	0	AAEL003347-PA OS=Aedes aegypti GN=AAEL003347 PE=4 SV=1
CCG026074.1	920	19	487	0.88741843	18.50508781	4.382163652	Up	7.73E-10	1.60E-09	AAEL006209-PA OS=Aedes aegypti GN=AAEL006209 PE=4 SV=1
CCG011676.1	1071	62	487	2.48751031	15.89606049	2.675894856	Up	0	0	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 SV=1

CCG019055.1	1938	107	487	2.37242601	8.784665009	1.888624179	Up	0	0	AAEL000811-PA (Fragment) OS=Aedes aegypti GN=GPRMTH7 PE=4 SV=2
CCG000103.1	711	153	487	9.24665175	23.94469872	1.372703323	Up	0	0	AAEL011150-PA OS=Aedes aegypti GN=AAEL011150 PE=4 SV=1
CCG008018.1	489	8	488	0.70298134	34.88678745	5.633050541	Up	2.19E-06	3.97E-06	AAEL000235-PA OS=Aedes aegypti GN=AAEL000235 PE=4 SV=1
CCG016427.1	7506	87	488	0.49805048	2.272800302	2.190107046	Up	0	0	Rapsynoid OS=Culex quinquefasciatus GN=CpipJ_CPIJ003545 PE=4 SV=1
CCG006558.2	1539	188	488	5.2490644	11.08488568	1.07846169	Up	1.80E-13	5.24E-13	AAEL009998-PA OS=Aedes aegypti GN=AAEL009998 PE=4 SV=1
CCG011166.1	2314	198	488	3.67675344	7.372359146	1.003693921	Up	0	0	AAEL002514-PA OS=Aedes aegypti GN=AAEL002514 PE=4 SV=1
CCG015431.1	49807	1095	488	0.94468367	0.342514889	-1.463664613	Down	8.61E-85	4.41E-84	AAEL003978-PA OS=Aedes aegypti GN=AAEL003978 PE=4 SV=1
CCG002365.1	891	55	489	2.65245275	19.1858556	2.854644145	Up	0	0	AAEL006068-PA OS=Aedes aegypti GN=AAEL006068 PE=4 SV=1
CCG018775.1	1478	64	489	1.8606651	11.56603338	2.636003859	Up	0	0	Ecdysteroid receptor OS=Aedes albopictus GN=EcR PE=2 SV=1
CCG022183.1	738	93	489	5.41488526	23.16341103	2.096845048	Up	4.04E-14	1.38E-13	Putative uncharacterized protein OS=Amblyomma maculatum PE=2 SV=1
CCG008945.1	1068	150	489	6.03507509	16.00617729	1.407185168	Up	0	0	AAEL010469-PA OS=Aedes aegypti GN=AAEL010469 PE=4 SV=1
CCG019825.2	1779	1372	489	33.1391095	9.609104745	-1.786060907	Down	9.06E-141	5.10E-140	AAEL009658-PA OS=Aedes aegypti GN=AAEL009658 PE=4 SV=1
CCG019135.1	2139	12	490	0.24106443	8.008207395	5.053988642	Up	1.30E-07	2.49E-07	AAEL002320-PA OS=Aedes aegypti GN=AAEL002320 PE=4 SV=1
CCG020448.1	1956	33	490	0.72494951	8.757441523	3.594557023	Up	1.31E-13	4.00E-13	AAEL003572-PA OS=Aedes aegypti GN=AAEL003572 PE=4 SV=1
CCG010446.1	1930	59	490	1.31358256	8.875417419	2.756308093	Up	0	0	AAEL007472-PA OS=Aedes aegypti GN=AAEL007472 PE=4 SV=1
CCG022159.1	966	134	490	5.96060501	17.73245923	1.572861952	Up	0	0	AAEL013557-PA OS=Aedes aegypti GN=AAEL013557 PE=4 SV=1

CCG009735.1	987	142	490	6.18206921	17.35517287	1.489204023	Up	2.62E-13	7.34E-13	AAEL010173-PA OS=Aedes aegypti GN=AAEL010173 PE=4 SV=1
CCG026283.1	1914	148	490	3.32263361	8.949611086	1.429497777	Up	0	0	AAEL013977-PA OS=Aedes aegypti GN=AAEL013977 PE=4 SV=1
CCG024382.1	1545	881	490	24.502483	11.08709102	-1.144047066	Down	3.31E-47	1.53E-46	AAEL004496-PA OS=Aedes aegypti GN=AAEL004496 PE=4 SV=1
CCG003111.1	477	9	491	0.81074971	35.98430586	5.471967417	Up	1.09E-06	2.01E-06	AAEL013145-PA OS=Aedes aegypti GN=AAEL013145 PE=4 SV=1
CCG012796.1	1643	72	491	1.88303158	10.44705654	2.471967417	Up	0	0	AAEL011868-PA OS=Aedes aegypti GN=AAEL011868 PE=4 SV=1
CCG018738.1	852	95	491	4.79122628	20.14614307	2.07203681	Up	0	0	AAEL010071-PA OS=Aedes aegypti GN=AAEL010071 PE=3 SV=1
CCG027068.1	2304	107	491	1.99555625	7.449875823	1.900425432	Up	0	0	AAEL000116-PA OS=Aedes aegypti GN=AAEL000116 PE=4 SV=1
CCG024291.1	4375	175	491	1.71878938	3.923317462	1.190681306	Up	3.55E-13	9.58E-13	AAEL013769-PA OS=Aedes aegypti GN=AAEL013769 PE=4 SV=1
CCG015834.2	1668	194	491	4.99767897	10.29047596	1.041979576	Up	5.97E-13	1.53E-12	AAEL005820-PA OS=Aedes aegypti GN=AAEL005820 PE=4 SV=1
CCG017973.1	1410	853	491	25.9951657	12.17341411	-1.094509513	Down	1.05E-42	4.72E-42	AAEL004119-PA OS=Aedes aegypti GN=AAEL004119 PE=4 SV=1
CCG016367.1	1416	40	492	1.21383431	12.1465199	3.322899614	Up	6.86E-14	2.24E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017880 PE=4 SV=1
CCG005954.1	1298	65	492	2.15179719	13.25074898	2.622459896	Up	0	0	U3 small nucleolar ribonucleoprotein component OS=Aedes aegypti PE=2 SV=1
CCG007785.1	546	120	492	9.44389772	31.50086478	1.737937114	Up	0	0	AAEL014722-PA OS=Aedes aegypti GN=AAEL014722 PE=4 SV=1
CCG008916.1	1221	129	492	4.53979997	14.0863818	1.633600454	Up	0	0	AAEL012140-PA OS=Aedes aegypti GN=AAEL012140 PE=4 SV=1
CCG027554.1	1897	162	492	3.66952926	9.066669569	1.304977706	Up	0	0	AAEL007472-PA OS=Aedes aegypti GN=AAEL007472 PE=4 SV=1
CCG023043.1	7149	938	492	5.63793692	2.405857067	-1.228616403	Down	5.30E-56	2.52E-55	AAEL001803-PA (Fragment) OS=Aedes aegypti GN=AAEL001803 PE=4 SV=1

CCG012694.1	1262	1106	492	37.6581034	13.62874182	-1.466307961	Down	7.47E-86	3.84E-85	AAEL009560-PA OS=Aedes aegypti GN=AAEL009560 PE=4 SV=1
CCG012547.1	843	19	493	0.96847563	20.44416423	4.399829527	Up	7.73E-10	1.60E-09	AAEL002806-PA OS=Aedes aegypti GN=AAEL002806 PE=4 SV=1
CCG004796.1	1971	46	493	1.00284515	8.744003272	3.124195084	Up	3.40E-14	1.18E-13	AAEL007344-PA OS=Aedes aegypti GN=AAEL007344 PE=4 SV=1
CCG006751.1	1110	50	493	1.93557363	15.52651392	3.00390085	Up	0	0	AAEL002267-PA OS=Aedes aegypti GN=AAEL002267 PE=4 SV=1
CCG011568.1	900	85	493	4.05825271	19.14936717	2.238366104	Up	0	0	AAEL001086-PA OS=Aedes aegypti GN=AAEL001086 PE=4 SV=1
CCG020987.1	1869	180	493	4.1383372	9.221204092	1.155903944	Up	3.32E-13	9.06E-13	AAEL006937-PA OS=Aedes aegypti GN=AAEL006937 PE=4 SV=1
CCG024861.1	1162	37	494	1.36822735	14.86178031	3.44122707	Up	5.17E-14	1.74E-13	-
CCG006998.1	1357	52	494	1.64659263	12.72615234	2.950240717	Up	0	0	AAEL014961-PA OS=Aedes aegypti GN=AAEL014961 PE=4 SV=1
CCG000039.1	1397	83	494	2.55296204	12.36176716	2.275641004	Up	1.32E-13	4.01E-13	AAEL006523-PA OS=Aedes aegypti GN=AAEL006523 PE=4 SV=1
CCG002614.1	946	192	494	8.72113007	18.25516779	1.065717935	Up	0	0	AAEL000964-PA OS=Aedes aegypti GN=AAEL000964 PE=4 SV=1
CCG008360.1	945	815	494	37.0585542	18.27448542	-1.019975814	Down	1.37E-36	5.99E-36	AAEL003989-PA OS=Aedes aegypti GN=AAEL003989 PE=4 SV=1
CCG014947.2	1197	34	495	1.22052713	14.45643024	3.566135078	Up	8.82E-14	2.79E-13	AAEL003654-PA OS=Aedes aegypti GN=AAEL003654 PE=4 SV=1
CCG004974.1	1370	68	495	2.13280435	12.63091022	2.566135078	Up	0	0	AAEL001644-PA OS=Aedes aegypti GN=AAEL001644 PE=4 SV=1
CCG020282.1	456	80	495	7.53854993	37.94812939	2.331669824	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000199 PE=4 SV=1
CCG002755.1	1409	170	495	5.18442504	12.28129667	1.244206983	Up	3.53E-13	9.52E-13	AAEL005727-PA OS=Aedes aegypti GN=AAEL005727 PE=4 SV=1
CCG021167.1	3636	1184	495	13.9923448	4.759171343	-1.555855447	Down	4.18E-100	2.21E-99	AAEL006355-PA OS=Aedes aegypti GN=SCRC1 PE=4 SV=2
CCG017484.1	375	9	496	1.03127363	46.23814741	5.486584513	Up	1.09E-06	2.01E-06	Putative salivary protein 34k-1 OS=Aedes albopictus PE=2 SV=1

CCG013375.1	1254	10	496	0.34266136	13.82719719	5.334581419	Up	5.43E-07	1.01E-06	AAEL005326-PA OS=Aedes aegypti GN=AAEL005326 PE=4 SV=1
CCG007831.1	885	25	496	1.21383431	19.59243534	4.012653324	Up	9.32E-12	2.08E-11	AAEL006107-PA OS=Aedes aegypti GN=AAEL006107 PE=4 SV=1
CCG000113.1	896	86	496	4.12432721	19.35190321	2.23024476	Up	0	0	AAEL000046-PA OS=Aedes aegypti GN=AAEL000046 PE=4 SV=1
CCG016555.1	2394	93	496	1.66925034	7.242817577	2.117350703	Up	4.04E-14	1.39E-13	AAEL004896-PA OS=Aedes aegypti GN=AAEL004896 PE=4 SV=1
CCG014201.1	4122	146	496	1.52197507	4.206527239	1.466684955	Up	9.33E-15	3.34E-14	AAEL004332-PA OS=Aedes aegypti GN=AAEL004332 PE=4 SV=1
CCG009495.1	1227	156	496	5.46314474	14.13146315	1.371107295	Up	1.42E-13	4.27E-13	Light protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010406 PE=4 SV=1
CCG004949.1	1677	43	497	1.10178807	10.36032412	3.233150491	Up	8.46E-14	2.70E-13	AAEL002048-PA OS=Aedes aegypti GN=AAEL002048 PE=3 SV=1
CCG013695.1	2286	96	497	1.80450329	7.600290269	2.074452745	Up	0	0	AAEL013594-PA OS=Aedes aegypti GN=AAEL013594 PE=4 SV=1
CCG019989.1	1256	103	497	3.52379193	13.83301239	1.972914718	Up	0	0	AAEL009269-PA OS=Aedes aegypti GN=AAEL009269 PE=4 SV=1
CCG016285.1	953	172	497	7.75529313	18.2311265	1.233150491	Up	1.35E-13	4.10E-13	AAEL000701-PA OS=Aedes aegypti GN=AAEL000701 PE=4 SV=1
CCG000587.1	2148	43	498	0.86019487	8.104851877	3.236050381	Up	8.46E-14	2.70E-13	AAEL006532-PA OS=Aedes aegypti GN=AAEL006532 PE=4 SV=1
CCG018918.1	12363	46	498	0.15988092	1.408171304	3.13875318	Up	3.40E-14	1.18E-13	Cytoplasmic dynein heavy chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ005887 PE=4 SV=1
CCG027738.1	485	80	498	7.08779128	35.89530275	2.340387041	Up	0	0	-
CCG004580.1	1839	111	498	2.59360551	9.466678539	1.86789927	Up	0	0	AAEL004708-PA (Fragment) OS=Aedes aegypti GN=AAEL004708 PE=4 SV=1
CCG000078.1	1242	111	498	3.84029029	14.01708682	1.86789927	Up	0	0	AAEL011326-PA OS=Aedes aegypti GN=GPRMTH5 PE=4 SV=2
CCG014210.1	516	144	498	11.9915538	33.738802	1.492390134	Up	7.53E-14	2.44E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010577 PE=4 SV=1

CCG024843.1	917	195	498	9.13751172	18.98497474	1.054984822	Up	2.26E-13	6.47E-13	AAEL010002-PA OS=Aedes aegypti GN=AAEL010002 PE=4 SV=1
CCG020785.1	2943	958	498	13.9874298	5.915467833	-1.24156671	Down	4.36E-58	2.09E-57	-
CCG005312.1	1902	3	499	0.06777561	9.171493223	7.080246708	Up	5.28E-05	8.85E-05	AAEL003088-PA OS=Aedes aegypti GN=AAEL003088 PE=4 SV=1
CCG010886.1	1627	107	499	2.82591371	10.72168415	1.923742223	Up	0	0	AAEL002298-PA OS=Aedes aegypti GN=AAEL002298 PE=4 SV=1
CCG017382.1	1073	118	499	4.72546942	16.2573906	1.78256616	Up	0	0	AAEL002790-PA OS=Aedes aegypti GN=AAEL002790 PE=4 SV=1
CCG007610.1	1317	163	499	5.31819798	13.24539112	1.316481055	Up	1.80E-13	5.23E-13	AAEL001978-PA (Fragment) OS=Aedes aegypti GN=AAEL001978 PE=4 SV=1
CCG028343.1	1438	167	499	4.99022648	12.13086238	1.281504917	Up	1.62E-13	4.78E-13	AAEL009116-PA OS=Aedes aegypti GN=AAEL009116 PE=4 SV=1
CCG010931.1	1069	179	499	7.19511927	16.31822274	1.181393432	Up	0	0	-
CCG014448.1	1962	1111	499	24.3319955	8.891019424	-1.452433892	Down	4.98E-85	2.56E-84	Voltage-gated calcium channel beta subunit OS=Anopheles gambiae PE=2 SV=1
CCG008211.1	382	1550	499	174.353635	45.66539296	-1.932843291	Down	7.44E-177	4.34E-176	-
CCG001904.1	1707	34	500	0.85587052	10.23968271	3.580634647	Up	8.82E-14	2.80E-13	AAEL012566-PA OS=Aedes aegypti GN=AAEL012566 PE=4 SV=1
CCG012151.1	1164	43	500	1.58736992	15.01644191	3.241832734	Up	8.46E-14	2.70E-13	AAEL009518-PA OS=Aedes aegypti GN=AAEL009518 PE=4 SV=1
CCG012582.1	1614	46	500	1.22466406	10.8297016	3.144535532	Up	3.40E-14	1.18E-13	AAEL003657-PA OS=Aedes aegypti GN=AAEL003657 PE=4 SV=1
CCG021938.1	1173	87	500	3.18701356	14.90122625	2.225153993	Up	0	0	AAEL002846-PA OS=Aedes aegypti GN=AAEL002846 PE=4 SV=1
CCG020091.1	1558	189	500	5.21263148	11.21895917	1.105855064	Up	2.34E-13	6.64E-13	AAEL001385-PA OS=Aedes aegypti GN=AAEL001385 PE=4 SV=1
CCG019821.1	1278	81	501	2.72343388	13.70430099	2.331129994	Up	4.46E-14	1.51E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_05553 PE=4 SV=1
CCG011299.1	1554	93	501	2.57154782	11.27033247	2.131821186	Up	4.04E-14	1.39E-13	AAEL010935-PA OS=Aedes aegypti GN=AAEL010935 PE=4 SV=1

CCG008654.1	969	127	501	5.63174024	18.07440316	1.68229531	Up	1.57E-13	4.67E-13	AAEL006039-PA OS=Aedes aegypti GN=AAEL006039 PE=3 SV=1
CCG010325.1	1413	4	502	0.12164115	12.41971333	6.673856758	Up	2.95E-05	5.04E-05	AAEL014380-PA OS=Aedes aegypti GN=AAEL014380 PE=4 SV=1
CCG005932.1	2385	156	502	2.81059899	7.358094314	1.388454539	Up	1.42E-13	4.26E-13	AAEL007215-PA OS=Aedes aegypti GN=AAEL007215 PE=4 SV=1
CCG003003.1	402	923	502	98.6593658	43.65436552	-1.17633008	Down	1.48E-51	6.96E-51	AAEL008672-PA OS=Aedes aegypti GN=AAEL008672 PE=3 SV=1
CCG025146.1	1979	111	503	2.4101266	8.885302282	1.882311927	Up	0	0	AAEL004853-PA OS=Aedes aegypti GN=AAEL004853 PE=3 SV=1
CCG005369.1	780	138	503	7.60233766	22.54360669	1.568203337	Up	1.49E-13	4.45E-13	AAEL009944-PA OS=Aedes aegypti GN=AAEL009944 PE=4 SV=1
CCG009789.1	1134	39	504	1.47779511	15.5370119	3.394190908	Up	7.33E-14	2.37E-13	AAEL008112-PA OS=Aedes aegypti GN=AAEL008112 PE=4 SV=1
CCG010951.1	453	98	504	9.29588078	38.89397681	2.064883283	Up	0	0	AAEL002550-PA OS=Aedes aegypti GN=AAEL002550 PE=4 SV=1
CCG008133.1	916	1144	504	53.6652581	19.23468504	-1.480278209	Down	5.46E-90	2.84E-89	-
CCG010908.1	710	3	505	0.18156226	24.86468982	7.097490281	Up	5.28E-05	8.85E-05	histone H1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ012377 PE=3 SV=1
CCG010219.1	1416	68	505	2.06351833	12.46746453	2.59498994	Up	0	0	AAEL010332-PA OS=Aedes aegypti GN=AAEL010332 PE=4 SV=1
CCG023361.1	999	123	505	5.29056793	17.67160137	1.739938276	Up	0	0	AAEL001688-PA OS=Aedes aegypti GN=AAEL001688 PE=4 SV=1
CCG016836.2	1443	167	505	4.97293533	12.23418556	1.298748489	Up	1.62E-13	4.77E-13	AAEL001859-PA OS=Aedes aegypti GN=AAEL001859 PE=4 SV=1
CCG022292.1	918	177	505	8.28501419	19.23086032	1.214847231	Up	0	0	AAEL009145-PA (Fragment) OS=Aedes aegypti GN=AAEL009145 PE=4 SV=1
CCG021977.1	285	180	505	27.1387798	61.94361323	1.190599685	Up	3.32E-13	9.06E-13	AAEL008928-PA OS=Aedes aegypti GN=AAEL008928 PE=3 SV=1
CCG028416.1	1220	1111	505	39.1306354	14.47043424	-1.43519032	Down	1.55E-83	7.94E-83	NADP+-dependent farnesol dehydrogenase 1 OS=Aedes aegypti GN=FOHSDR-1 PE=2 SV=1

CCG013485.1	1884	1637	505	37.3362291	9.370451046	-1.994385825	Down	1.23E-194	7.29E-194	AAEL013821-PA OS=Aedes aegypti GN=AAEL013821 PE=4 SV=1
CCG018683.1	1260	40	506	1.36411856	14.03880004	3.363378684	Up	6.86E-14	2.25E-13	AAEL004837-PA OS=Aedes aegypti GN=AAEL004837 PE=4 SV=1
CCG012898.1	1359	61	506	1.92873717	13.01610599	2.754569441	Up	1.75E-14	6.17E-14	AAEL006220-PA OS=Aedes aegypti GN=AAEL006220 PE=4 SV=1
CCG024456.1	821	80	506	4.18706306	21.54553964	2.363378684	Up	0	0	Exosomal 3'-5' exoribonuclease complex subunit ski4 (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG017591.1	750	117	506	6.7032786	23.58518406	1.814942059	Up	9.44E-14	2.97E-13	AAEL012195-PA OS=Aedes aegypti GN=AAEL012195 PE=4 SV=1
CCG016255.1	417	122	506	12.5714811	42.4193958	1.754569441	Up	8.04E-14	2.58E-13	AAEL011959-PA OS=Aedes aegypti GN=AAEL011959 PE=4 SV=1
CCG020985.1	363	174	506	20.5970629	48.72971914	1.242363283	Up	1.78E-13	5.20E-13	AAEL001341-PA OS=Aedes aegypti GN=AAEL001341 PE=4 SV=1
CCG011451.2	1672	192	506	4.93432359	10.5794785	1.100344278	Up	0	0	AAEL012479-PA OS=Aedes aegypti GN=AAEL012479 PE=4 SV=1
CCG004582.2	1782	878	506	21.1713956	9.926424269	-1.092770351	Down	8.47E-44	3.84E-43	Uncharacterized protein OS=Anopheles darlingi GN=AND_22059 PE=4 SV=1
CCG010506.1	1344	63	507	2.01420631	13.18738566	2.710875217	Up	0	0	AAEL010138-PA OS=Aedes aegypti GN=AAEL010138 PE=4 SV=1
CCG017936.2	720	93	507	5.55025739	24.61645323	2.14899633	Up	4.04E-14	1.38E-13	-
CCG016974.1	639	163	507	10.9609808	27.73684871	1.339426987	Up	1.80E-13	5.24E-13	AAEL011217-PA OS=Aedes aegypti GN=AAEL011217 PE=4 SV=1
CCG010621.1	480	182	507	16.292691	36.92467984	1.180360501	Up	0	0	AAEL001959-PA OS=Aedes aegypti GN=AAEL001959 PE=4 SV=1
CCG027818.1	1560	188	507	5.17840391	11.36143995	1.133566289	Up	1.80E-13	5.24E-13	AAEL008100-PA OS=Aedes aegypti GN=AAEL008100 PE=4 SV=1
CCG020793.1	730	1038	507	61.0994309	24.27924154	-1.331435588	Down	7.47E-70	3.70E-69	AAEL013262-PA OS=Aedes aegypti GN=AAEL013262 PE=4 SV=1
CCG013176.1	830	1726	507	89.3563397	21.35403172	-2.065061608	Down	1.03E-214	6.21E-214	AAEL000657-PA OS=Aedes aegypti GN=AAEL000657 PE=4 SV=1
CCG019995.1	1413	100	508	3.04102863	12.56815612	2.047141701	Up	0	0	AAEL009259-PA OS=Aedes aegypti GN=AAEL009259 PE=4 SV=1

CCG009137.1	2251	127	508	2.42432532	7.889295691	1.702313204	Up	1.57E-13	4.66E-13	AAEL005631-PA OS=Aedes aegypti GN=AAEL005631 PE=4 SV=1
CCG018625.1	2175	143	508	2.82513657	8.164967632	1.531126554	Up	0	0	AAEL002016-PA OS=Aedes aegypti GN=AAEL002016 PE=4 SV=1
CCG018945.1	1844	2398	508	55.879297	9.630588178	-2.536618053	Down	0	0	Cytokrome P450 OS=Aedes albopictus GN=CYP6Y7 PE=2 SV=1
CCG010482.1	786	0	509	0.001	22.63837516	14.46648279	Up	4.44E-16	1.65E-15	AAEL013288-PA OS=Aedes aegypti GN=AAEL013288 PE=4 SV=1
CCG024402.1	1734	43	509	1.06557012	10.26168563	3.267570295	Up	8.46E-14	2.70E-13	AAEL002751-PA OS=Aedes aegypti GN=AAEL002751 PE=4 SV=1
CCG016219.1	551	75	509	5.84887495	32.29358054	2.465016359	Up	0	0	AAEL010121-PA OS=Aedes aegypti GN=AAEL010121 PE=4 SV=1
CCG023251.1	1545	99	509	2.75340047	11.51699863	2.06447843	Up	0	0	AAEL010835-PA (Fragment) OS=Aedes aegypti GN=AAEL010835 PE=4 SV=1
CCG015296.1	930	105	509	4.85142165	19.13307836	1.979589532	Up	0	0	AAEL004797-PA OS=Aedes aegypti GN=AAEL004797 PE=4 SV=1
CCG028068.1	1233	133	509	4.63501598	14.43127565	1.638552614	Up	9.90E-14	3.11E-13	AAEL001740-PA OS=Aedes aegypti GN=AAEL001740 PE=4 SV=1
CCG003900.1	2046	137	509	2.87725007	8.696853801	1.595802967	Up	2.90E-13	8.03E-13	AAEL013092-PA OS=Aedes aegypti GN=AAEL013092 PE=4 SV=1
CCG024123.2	1515	977	509	27.7105153	11.745058	-1.238379702	Down	5.71E-59	2.74E-58	Cytokrome P450 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005957 PE=3 SV=1
CCG015550.1	1123	1581	509	60.4943459	15.84484673	-1.932786602	Down	2.41E-180	1.41E-179	AAEL003019-PA OS=Aedes aegypti GN=AAEL003019 PE=4 SV=1
CCG006606.1	1653	67	510	1.74166498	10.78567523	2.63057745	Up	0	0	AAEL001175-PA OS=Aedes aegypti GN=AAEL001175 PE=4 SV=1
CCG001963.1	528	83	510	6.75471207	33.76651734	2.321627209	Up	1.32E-13	4.02E-13	AAEL006781-PA OS=Aedes aegypti GN=AAEL006781 PE=4 SV=1
CCG021795.1	849	95	510	4.80815641	20.99967156	2.126811032	Up	0	0	AAEL005939-PA OS=Aedes aegypti GN=AAEL005939 PE=4 SV=1
CCG026106.1	1676	97	510	2.48691185	10.63766179	2.096753798	Up	3.62E-14	1.25E-13	AAEL007649-PA OS=Aedes aegypti GN=AAEL007649 PE=4 SV=1

CCG025632.1	5337	1024	510	8.24452094	3.340588562	-1.303333359	Down	9.68E-67	4.75E-66	AAEL006809-PA OS=Aedes aegypti GN=AAEL006809 PE=4 SV=1
CCG004225.1	1383	54	511	1.677777706	12.91661564	2.944605183	Up	0	0	Serine/threonine-protein kinase RIO1 OS=Aedes aegypti GN=AAEL011857 PE=3 SV=1
CCG005538.1	568	72	511	5.44686777	31.45013984	2.529567683	Up	0	0	Odorant-binding protein 2 OS=Aedes aegypti PE=2 SV=1
CCG015160.1	2555	169	511	2.84222511	6.991655354	1.298613248	Up	0	0	AAEL005472-PA OS=Aedes aegypti GN=AAEL005472 PE=4 SV=1
CCG015250.1	2053	20	512	0.41860433	8.718284319	4.380385109	Up	3.69E-10	7.72E-10	AAEL003542-PA OS=Aedes aegypti GN=AAEL003542 PE=4 SV=1
CCG013648.1	1341	48	512	1.53806656	13.3472317	3.117350703	Up	5.35E-14	1.79E-13	AAEL009971-PA OS=Aedes aegypti GN=AAEL009971 PE=4 SV=1
CCG009494.1	1719	71	512	1.77478252	10.41223834	2.552566084	Up	0	0	AAEL001152-PA OS=Aedes aegypti GN=AAEL001152 PE=4 SV=1
CCG010830.1	665	109	512	7.04315951	26.91524467	1.934128879	Up	0	0	AAEL001264-PA OS=Aedes aegypti GN=AAEL001264 PE=4 SV=1
CCG019633.1	870	120	512	5.92685995	20.57314679	1.795422608	Up	0	0	AAEL014038-PA OS=Aedes aegypti GN=AAEL014038 PE=3 SV=1
CCG027023.1	666	129	512	8.32296661	26.87483139	1.691085948	Up	0	0	AAEL013320-PA OS=Aedes aegypti GN=AAEL013320 PE=4 SV=1
CCG021792.1	1362	130	512	4.10136968	13.14143738	1.679945391	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000820 PE=4 SV=1
CCG021571.1	2481	1650	512	28.5772117	7.214283639	-1.985937105	Down	4.57E-195	2.72E-194	AAEL004074-PA OS=Aedes aegypti GN=AAEL004074 PE=4 SV=1
CCG002690.1	1260	7	513	0.23872075	14.23301269	5.897773297	Up	4.31E-06	7.71E-06	AAEL002556-PA (Fragment) OS=Aedes aegypti GN=AAEL002556 PE=4 SV=1
CCG025158.1	1743	19	513	0.46840216	10.28892483	4.457200706	Up	7.73E-10	1.60E-09	AAEL011085-PA OS=Aedes aegypti GN=AAEL011085 PE=4 SV=1
CCG003093.1	990	39	513	1.69274712	18.11474342	3.419726001	Up	7.33E-14	2.38E-13	AAEL000157-PA OS=Aedes aegypti GN=AAEL000157 PE=4 SV=1

CCG020673.1	1302	57	513	1.8811635	13.77388324	2.872238205	Up	0	0	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001406 PE=4 SV=1
CCG017941.1	855	71	513	3.56824697	20.97496606	2.5553811	Up	0	0	AAEL005424-PA OS=Aedes aegypti GN=AAEL005424 PE=4 SV=1
CCG024388.1	1734	99	513	2.45328935	10.34232756	2.075771599	Up	0	0	AAEL002751-PA OS=Aedes aegypti GN=AAEL002751 PE=4 SV=1
CCG011086.1	363	120	513	14.2048709	49.40384569	1.798237624	Up	0	0	AAEL013060-PA OS=Aedes aegypti GN=AAEL013060 PE=4 SV=1
CCG007317.1	738	160	513	9.31593162	24.30026556	1.383200125	Up	4.24E-14	1.44E-13	AAEL009615-PA OS=Aedes aegypti GN=GPROP8 PE=2 SV=1
CCG021774.1	1374	38	514	1.1883915	13.07755041	3.46001024	Up	4.26E-14	1.45E-13	AAEL011655-PA OS=Aedes aegypti GN=AAEL011655 PE=3 SV=1
CCG017810.1	1851	112	514	2.60000555	9.707484744	1.900582831	Up	0	0	AAEL011465-PA OS=Aedes aegypti GN=AAEL011465 PE=4 SV=1
CCG028137.1	1663	118	514	3.04896493	10.80490334	1.825294704	Up	0	0	AAEL012146-PA (Fragment) OS=Aedes aegypti GN=AAEL012146 PE=4 SV=1
CCG026922.1	558	175	514	13.4761712	32.20171015	1.256726641	Up	3.55E-13	9.58E-13	AAEL009085-PA OS=Aedes aegypti GN=AAEL009085 PE=4 SV=1
CCG006273.1	2284	201	514	3.78148715	7.867142846	1.056886062	Up	0	0	AAEL011074-PA OS=Aedes aegypti GN=AAEL011074 PE=4 SV=1
CCG012308.1	1404	203	514	6.21286049	12.79811557	1.042601836	Up	9.28E-14	2.93E-13	AAEL010223-PA OS=Aedes aegypti GN=AAEL010223 PE=4 SV=1
CCG010872.1	2364	1776	514	32.2818311	7.600911278	-2.086478113	Down	7.41E-224	4.49E-223	AAEL012094-PA OS=Aedes aegypti GN=AAEL012094 PE=4 SV=1
CCG019339.1	1851	1936	514	44.9429531	9.707484744	-2.210925484	Down	1.38E-262	8.53E-262	AGAP001797-PA OS=Anopheles gambiae GN=AgaP_AGAP001797 PE=3 SV=4
CCG006896.1	1464	2891	514	84.8534855	12.27360264	-2.78941514	Down	0	0	AAEL011981-PA OS=Aedes aegypti GN=AAEL011981 PE=3 SV=1
CCG021728.1	2217	89	515	1.7249916	8.120664203	2.235008395	Up	0	0	AAEL012059-PA OS=Aedes aegypti GN=AAEL012059 PE=4 SV=1
CCG028465.1	2151	102	515	2.03761643	8.369833816	2.038316484	Up	0	0	AAEL007067-PA OS=Aedes aegypti GN=AAEL007067 PE=4 SV=1

CCG021365.1	807	121	515	6.44279788	22.3091853	1.791878589	Up	0	0	AGAP005928-PA OS=Anopheles gambiae GN=AGAP005928 PE=4 SV=1
CCG011335.1	1611	1413	515	37.6885382	11.1753647	-1.753803924	Down	2.73E-141	1.54E-140	AAEL010294-PA OS=Aedes aegypti GN=AAEL010294 PE=4 SV=1
CCG013128.1	615	2411	515	168.455334	29.27400413	-2.52467411	Down	0	0	AAEL006439-PA OS=Aedes aegypti GN=AAEL006439 PE=4 SV=1
CCG006253.1	318	92	516	12.4314955	56.72475099	2.189978503	Up	1.44E-14	5.11E-14	AAEL010581-PA OS=Aedes aegypti GN=AAEL010581 PE=4 SV=1
CCG013389.1	1624	162	516	4.28638978	11.10743277	1.373690456	Up	0	0	AAEL002489 OS=Aedes aegypti GN=AAEL002489 PE=2 SV=1
CCG026123.1	402	0	517	0.001	44.95877883	15.45631523	Up	4.44E-16	1.63E-15	AAEL013734-PA OS=Aedes aegypti GN=AAEL013734 PE=4 SV=1
CCG014017.1	1242	103	517	3.56351261	14.55187527	2.029833147	Up	0	0	AAEL004420-PA OS=Aedes aegypti GN=AAEL004420 PE=4 SV=1
CCG015503.1	1221	114	517	4.01191625	14.80215323	1.88344366	Up	8.48E-14	2.70E-13	AAEL012838-PA OS=Aedes aegypti GN=AAEL012838 PE=4 SV=1
CCG024790.1	1782	120	517	2.89358482	10.1422161	1.809443079	Up	0	0	AAEL001711-PA OS=Aedes aegypti GN=AAEL001711 PE=3 SV=1
CCG019667.1	1280	140	517	4.69981472	14.11986648	1.587050657	Up	0	0	AAEL000724-PA OS=Aedes aegypti GN=AAEL000724 PE=4 SV=1
CCG013644.1	507	159	517	13.4757156	35.64778913	1.403450719	Up	5.53E-14	1.84E-13	AAEL009957-PA OS=Aedes aegypti GN=AAEL009957 PE=4 SV=1
CCG011161.1	1068	197	517	7.92606528	16.92268642	1.094281855	Up	0	0	AAEL006611-PA OS=Aedes aegypti GN=AAEL006611 PE=4 SV=1
CCG022509.1	879	31	518	1.51542864	20.60112329	3.764925181	Up	2.40E-13	6.76E-13	AAEL013852-PA OS=Aedes aegypti GN=AAEL013852 PE=4 SV=1
CCG013853.1	1817	32	518	0.75675922	9.966091012	3.719121492	Up	1.77E-13	5.17E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008808 PE=4 SV=1
CCG004804.1	2472	63	518	1.09510246	7.325399421	2.741841568	Up	0	0	AAEL007237-PA OS=Aedes aegypti GN=AAEL007237 PE=4 SV=1
CCG000122.1	3033	76	518	1.07672266	5.970454127	2.471193978	Up	0	0	histone transcription regulator OS=Culex quinquefasciatus GN=CpipJ_CPIJ002429 PE=4 SV=1

CCG024507.1	1758	148	518	3.61747481	10.30056164	1.509668126	Up	0	0	AAEL012329-PA (Fragment) OS=Aedes aegypti GN=AAEL012329 PE=3 SV=1
CCG006576.1	1668	60	519	1.54567391	10.87730554	2.815013337	Up	0	0	AAEL012469-PA OS=Aedes aegypti GN=AAEL012469 PE=4 SV=1
CCG022463.1	749	88	519	5.04851355	24.22342543	2.262472314	Up	2.93E-14	1.02E-13	AAEL008891-PA OS=Aedes aegypti GN=AAEL008891 PE=4 SV=1
CCG026617.1	1361	92	519	2.9046404	13.3308932	2.198341976	Up	1.44E-14	5.11E-14	AAEL004740-PA OS=Aedes aegypti GN=AAEL004740 PE=4 SV=1
CCG000639.1	732	96	519	5.63537503	24.78599132	2.136941431	Up	0	0	AGAP003872-PA OS=Anopheles gambiae GN=AgaP_AGAP003872 PE=4 SV=4
CCG012142.1	2688	83	520	1.32681844	6.762761876	2.349641585	Up	1.32E-13	4.01E-13	AAEL011963-PA OS=Aedes aegypti GN=AAEL011963 PE=4 SV=1
CCG004227.1	1531	104	520	2.91891078	11.87348395	2.024241299	Up	0	0	AAEL015347-PA OS=Aedes aegypti GN=AAEL015347 PE=4 SV=1
CCG002638.1	1251	112	520	3.84701061	14.53101832	1.917326095	Up	0	0	AAEL014440-PA OS=Aedes aegypti GN=AAEL014440 PE=4 SV=1
CCG023699.1	2965	131	520	1.89849418	6.130962537	1.691258015	Up	1.26E-13	3.87E-13	AAEL003447-PA OS=Aedes aegypti GN=AAEL003447 PE=4 SV=1
CCG016645.1	1260	163	520	5.55878313	14.42722533	1.375952863	Up	1.80E-13	5.25E-13	AAEL005729-PA OS=Aedes aegypti GN=HOT_AEDAE PE=4 SV=1
CCG011657.1	1008	210	520	8.95202804	18.03403167	1.010435499	Up	0	0	AAEL000394-PA OS=Aedes aegypti GN=AAEL000394 PE=4 SV=1
CCG014125.1	459	996	520	93.2415156	39.60414798	-1.235320915	Down	7.46E-60	3.59E-59	AAEL008671-PA OS=Aedes aegypti GN=AAEL008671 PE=4 SV=1
CCG009201.1	930	69	521	3.18807708	19.5841529	2.618928309	Up	0	0	AAEL011851-PA OS=Aedes aegypti GN=AAEL011851 PE=4 SV=1
CCG000912.1	1282	84	521	2.81548963	14.20691279	2.335135343	Up	0	0	AAEL013048-PA OS=Aedes aegypti GN=AAEL013048 PE=4 SV=1
CCG015696.1	1605	1630	521	43.6390451	11.34782691	-1.943203483	Down	3.23E-187	1.91E-186	AAEL014318-PA OS=Aedes aegypti GN=AAEL014318 PE=4 SV=1
CCG022692.1	16935	2972	521	7.54095372	1.075480496	-2.809765634	Down	0	0	AAEL005244-PA OS=Aedes aegypti GN=AAEL005244 PE=4 SV=1

CCG012955.1	933	8720	521	401.603522	19.52118135	-4.362659654	Down	0	0	AAEL006586-PA (Fragment) OS=Aedes aegypti GN=AAEL006586 PE=3 SV=1
CCG012842.1	702	47	522	2.87689106	25.99461606	3.175630349	Up	0	0	AAEL013274-PA OS=Aedes aegypti GN=AAEL013274 PE=3 SV=1
CCG027396.1	1549	52	522	1.44249593	11.78064588	3.029779482	Up	0	0	AAEL000839-PA OS=Aedes aegypti GN=AAEL000839 PE=4 SV=1
CCG000863.1	654	110	522	7.22732539	27.90247779	1.948859487	Up	0	0	-
CCG016794.1	2251	181	522	3.45514081	8.106717226	1.230373313	Up	1.53E-13	4.55E-13	AAEL011370-PA OS=Aedes aegypti GN=AAEL011370 PE=4 SV=1
CCG001649.1	402	1813	523	193.791365	45.48054416	-2.09118287	Down	3.81E-229	2.32E-228	Cytochrome b-c1 complex subunit 7 OS=Aedes aegypti GN=AAEL007868 PE=3 SV=1
CCG002039.1	1387	62	524	1.92078122	13.20702021	2.781539895	Up	0	0	AAEL013760-PA OS=Aedes aegypti GN=AAEL013760 PE=3 SV=1
CCG022758.1	1190	76	524	2.74428557	15.39339246	2.487808692	Up	0	0	AAEL002010-PA OS=Aedes aegypti GN=AAEL002010 PE=4 SV=1
CCG004061.1	1290	99	524	3.29767731	14.20010622	2.106379585	Up	0	0	AAEL012841-PA OS=Aedes aegypti GN=AAEL012841 PE=4 SV=1
CCG027082.1	450	109	524	10.4082246	40.70697117	1.967551881	Up	0	0	-
CCG010175.1	1263	171	524	5.81775504	14.50367144	1.31788369	Up	3.42E-13	9.25E-13	Transcription initiation factor TFIID subunit 7 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008789 PE=4 SV=1
CCG014382.1	2185	182	524	3.5791724	8.383586741	1.227941565	Up	0	0	AAEL012288-PA OS=Aedes aegypti GN=AAEL012288 PE=3 SV=1
CCG027551.3	1874	1384	524	31.7343184	9.774886355	-1.698892022	Down	2.20E-132	1.23E-131	Methyltetrahydropteroyl reductase OS=Culex quinquefasciatus GN=CpipJ_CPIJ003297 PE=2 SV=1
CCG013191.1	396	4135	524	448.686497	46.25792179	-3.277935409	Down	0	0	AAEL002606-PA OS=Aedes aegypti GN=AAEL002606 PE=4 SV=1
CCG020171.1	1808	0	525	0.001	10.15104829	13.3093411	Up	4.44E-16	1.64E-15	AAEL003429-PA OS=Aedes aegypti GN=AAEL003429 PE=4 SV=1
CCG007387.1	875	60	525	2.94649609	20.97496606	2.831596221	Up	0	0	AAEL007655-PA OS=Aedes aegypti GN=AAEL007655 PE=4 SV=1

CCG027081.1	1503	82	525	2.34432351	12.21097492	2.380934812	Up	0	0	AAEL007454-PA OS=Aedes aegypti GN=AAEL007454 PE=4 SV=1
CCG011083.1	944	128	525	5.82640469	19.44183825	1.738486816	Up	4.15E-14	1.42E-13	AAEL013998-PA OS=Aedes aegypti GN=AAEL013998 PE=4 SV=1
CCG012332.1	483	164	525	14.5901376	37.99812693	1.380934812	Up	1.14E-13	3.54E-13	AAEL012723-PA OS=Aedes aegypti GN=AAEL012723 PE=4 SV=1
CCG024326.1	3354	41	526	0.52527106	5.482425039	3.383680188	Up	1.02E-13	3.19E-13	AAEL008178-PA OS=Aedes aegypti GN=AAEL008178 PE=3 SV=1
CCG008235.1	1179	50	526	1.82229579	15.59631347	3.097376003	Up	0	0	AAEL006286-PA OS=Aedes aegypti GN=AAEL006286 PE=4 SV=1
CCG019390.1	1059	72	526	2.92145504	17.36360112	2.571307192	Up	0	0	AAEL007356-PA OS=Aedes aegypti GN=AAEL007356 PE=4 SV=1
CCG014258.1	435	92	526	9.08785192	42.27138755	2.217670237	Up	1.44E-14	5.12E-14	AAEL010680-PA OS=Aedes aegypti GN=AAEL010680 PE=4 SV=1
CCG003345.1	1212	126	526	4.46715063	15.17166137	1.76395227	Up	1.34E-13	4.06E-13	AAEL005564-PA OS=Aedes aegypti GN=AAEL005564 PE=4 SV=1
CCG013537.1	1860	139	526	3.21117909	9.886050313	1.62229112	Up	1.46E-13	4.37E-13	AAEL002478-PA OS=Aedes aegypti GN=AAEL002478 PE=4 SV=1
CCG002495.1	1173	177	526	6.48392415	15.67609001	1.273626643	Up	0	0	AAEL000044-PB OS=Aedes aegypti GN=AAEL000044 PE=3 SV=1
CCG015843.1	4476	1450	526	13.9200436	4.108144232	-1.760604992	Down	1.01E-145	5.70E-145	Uncharacterized protein OS=Anopheles darlingi GN=AND_01925 PE=4 SV=1
CCG003547.2	12095	1566	526	5.56350594	1.520302074	-1.871636304	Down	4.71E-171	2.73E-170	-
CCG018851.1	1881	151	527	3.4494577	9.794264678	1.505567616	Up	0	0	Acyl-coenzyme A oxidase OS=Aedes aegypti GN=AAEL000370 PE=3 SV=1
CCG012581.2	1623	42	528	1.11197095	11.37274808	3.3543899	Up	7.66E-14	2.47E-13	AAEL003679-PA OS=Aedes aegypti GN=AAEL003679 PE=4 SV=1
CCG001898.1	2796	69	528	1.06041191	6.601562996	2.638182866	Up	0	0	AAEL010387-PA OS=Aedes aegypti GN=AAEL010387 PE=4 SV=1
CCG022441.1	1482	71	528	2.05860402	12.45477067	2.596960204	Up	0	0	AAEL004007-PA OS=Aedes aegypti GN=AAEL004007 PE=3 SV=1
CCG017419.1	1770	173	528	4.19986672	10.42823172	1.312079096	Up	3.60E-13	9.69E-13	AAEL012076-PA OS=Aedes aegypti GN=AAEL012076 PE=4 SV=1

CCG019362.1	13426	1405	528	4.49668383	1.374792949	-1.709647092	Down	1.45E-135	8.12E-135	AAEL005179-PA OS=Aedes aegypti GN=AAEL005179 PE=4 SV=1
CCG018673.1	1630	52	529	1.37081362	11.34535485	3.048997398	Up	0	0	AAEL009727-PA OS=Aedes aegypti GN=AAEL009727 PE=4 SV=1
CCG007096.1	935	65	529	2.9872008	19.77853306	2.727069303	Up	0	0	AAEL011851-PA OS=Aedes aegypti GN=AAEL011851 PE=4 SV=1
CCG008505.1	2531	129	529	2.1900813	7.306569898	1.738209861	Up	0	0	AAEL014815-PA (Fragment) OS=Aedes aegypti GN=AAEL014815 PE=4 SV=1
CCG007928.1	687	151	529	9.44458505	26.91838197	1.511032377	Up	0	0	AAEL010931-PA OS=Aedes aegypti GN=AAEL010931 PE=2 SV=1
CCG016673.1	1143	163	529	6.12779242	16.17928995	1.400708962	Up	1.80E-13	5.24E-13	AAEL000459-PA OS=Aedes aegypti GN=AAEL000459 PE=4 SV=1
CCG025467.2	664	1165	529	75.3911759	27.8507958	-1.436677124	Down	1.22E-87	6.31E-87	AAEL009810-PA OS=Aedes aegypti GN=AAEL009810 PE=2 SV=1
CCG010991.1	2460	53	530	0.92577071	7.531661256	3.024241299	Up	0	0	AAEL013940-PA OS=Aedes aegypti GN=AAEL013940 PE=4 SV=1
CCG017306.1	639	107	530	7.19524508	28.99512784	2.010694767	Up	0	0	AAEL006599-PA OS=Aedes aegypti GN=AAEL006599 PE=4 SV=1
CCG013005.1	1152	1071	530	39.9484251	16.08323497	-1.312581011	Down	1.79E-70	8.87E-70	AAEL011393-PA OS=Aedes aegypti GN=AAEL011393 PE=4 SV=1
CCG020516.1	1472	48	531	1.401187	12.61062837	3.169918754	Up	5.35E-14	1.79E-13	AAEL014536-PA OS=Aedes aegypti GN=AAEL014536 PE=4 SV=1
CCG025793.1	519	53	531	4.38804612	35.76656063	3.0269608	Up	0	0	AAEL011330-PA OS=Aedes aegypti GN=AAEL011330 PE=4 SV=1
CCG001252.1	1200	67	531	2.39914352	15.46903747	2.688792064	Up	0	0	AAEL006915-PA OS=Aedes aegypti GN=AAEL006915 PE=4 SV=1
CCG005714.1	1018	163	531	6.88022273	18.23462177	1.4061531	Up	1.80E-13	5.24E-13	AAEL003206-PA OS=Aedes aegypti GN=AAEL003206 PE=4 SV=1
CCG011401.1	2379	181	531	3.26924	7.802793176	1.255035368	Up	1.53E-13	4.55E-13	AAEL015283-PA OS=Aedes aegypti GN=AAEL015283 PE=4 SV=1
CCG014041.1	1161	207	531	7.66127051	15.98866922	1.061394297	Up	6.88E-15	2.47E-14	AAEL002078-PA OS=Aedes aegypti GN=AAEL002078 PE=4 SV=1

CCG016111.1	6177	2432	531	16.9179852	3.00515541	-2.493046259	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006967 PE=3 SV=1
CCG003532.1	1764	150	532	3.653889	10.54297236	1.528776949	Up	0	0	AAEL009340-PA OS=Aedes aegypti GN=AAEL009340 PE=4 SV=1
CCG007611.1	2388	23	533	0.4138626	7.802663953	4.236742971	Up	4.01E-11	8.67E-11	AAEL004302-PA OS=Aedes aegypti GN=AAEL004302 PE=4 SV=1
CCG022222.1	1026	102	533	4.27184496	18.16058628	2.087879585	Up	0	0	AAEL014296-PA OS=Aedes aegypti GN=AAEL014296 PE=4 SV=1
CCG004643.3	5330	140	533	1.12866095	3.495827677	1.63102191	Up	0	0	Zinc finger protein 7 OS=Mus musculus GN=Zfp7 PE=2 SV=1
CCG014165.1	2065	162	533	3.37099129	9.023129065	1.420454924	Up	0	0	AAEL002821-PA OS=Aedes aegypti GN=AAEL002821 PE=4 SV=1
CCG007770.1	1099	70	534	2.73692577	16.98609627	2.633726119	Up	0	0	AAEL009501-PA OS=Aedes aegypti GN=AAEL009501 PE=3 SV=1
CCG000412.1	466	120	534	11.0651677	40.05948454	1.85611854	Up	0	0	-
CCG000138.1	1215	176	534	6.22442246	15.36437843	1.303577517	Up	0	0	AAEL009984-PA OS=Aedes aegypti GN=AAEL009984 PE=4 SV=1
CCG017729.1	7689	1125	534	6.28702711	2.427847548	-1.372700151	Down	4.06E-79	2.06E-78	AAEL011164-PA (Fragment) OS=Aedes aegypti GN=AAEL011164 PE=4 SV=1
CCG014916.1	393	76	535	8.30966878	47.58951164	2.517780772	Up	0	0	AAEL006346-PA OS=Aedes aegypti GN=AAEL006346 PE=4 SV=1
CCG016164.1	1245	104	535	3.58943968	15.02223138	2.065268567	Up	0	0	AAEL012034-PA OS=Aedes aegypti GN=AAEL012034 PE=4 SV=1
CCG001063.1	1809	198	535	4.70315503	10.33868329	1.136351665	Up	0	0	AAEL010761-PA OS=Aedes aegypti GN=AAEL010761 PE=4 SV=1
CCG004833.1	591	35	536	2.54473894	31.70496844	3.639119377	Up	6.39E-14	2.10E-13	AAEL010109-PA OS=Aedes aegypti GN=AAEL010109 PE=4 SV=1
CCG009904.1	657	114	536	7.45593569	28.51999444	1.93551238	Up	8.48E-14	2.70E-13	AAEL009051-PA (Fragment) OS=Aedes aegypti GN=TPX5 PE=4 SV=2
CCG024074.1	2973	141	536	2.0379188	6.302602203	1.628851042	Up	0	0	AAEL005848-PA OS=Aedes aegypti GN=AAEL005848 PE=4 SV=1

CCG002692.1	4035	2126	536	22.6403112	4.643776047	-2.285523487	Down	2.41E-300	1.52E-299	Bruchpilot OS=Culex quinquefasciatus GN=CpipJ_CPIJ017714 PE=4 SV_1
CCG026180.1	583	7	537	0.51593163	32.19999078	5.96373656	Up	4.31E-06	7.69E-06	-
CCG027212.1	1176	64	537	2.33848896	15.96309067	2.771091482	Up	0	0	Ferrochelatase OS=Aedes aegypti GN=AAEL005415 PE=3 SV=1
CCG000366.1	2021	80	537	1.70092962	9.288765278	2.449163387	Up	0	0	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV_1
CCG016434.1	946	95	537	4.31514248	19.84418037	2.201235873	Up	0	0	AAEL003650-PA OS=Aedes aegypti GN=AAEL003650 PE=4 SV_1
CCG009661.1	909	139	537	6.57072949	20.65191928	1.652150409	Up	1.46E-13	4.37E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001384 PE=4 SV_1
CCG005110.1	1854	42	538	0.97342441	10.14431117	3.381458144	Up	7.66E-14	2.47E-13	AAEL001175-PA OS=Aedes aegypti GN=AAEL001175 PE=4 SV_1
CCG017765.1	1017	140	538	5.91520437	18.49316903	1.644492549	Up	0	0	AAEL006221-PA OS=Aedes aegypti GN=AAEL006221 PE=4 SV_1
CCG002288.1	1422	16	539	0.48348506	13.2507111	4.776454667	Up	7.09E-09	1.42E-08	AAEL004468-PA OS=Aedes aegypti GN=AAEL004468 PE=4 SV_1
CCG000468.1	457	142	539	13.3516462	41.23087786	1.626707547	Up	2.62E-13	7.34E-13	AAEL012284-PA OS=Aedes aegypti GN=AAEL012284 PE=4 SV_1
CCG000351.1	978	69	540	3.03160704	19.30211601	2.670604344	Up	0	0	AAEL004625-PA OS=Aedes aegypti GN=AAEL004625 PE=4 SV_1
CCG012351.1	351	100	540	12.2420896	53.78196427	2.135272611	Up	0	0	AAEL000456-PA OS=Aedes aegypti GN=AAEL000456 PE=4 SV_1
CCG004341.1	2604	125	540	2.06267927	7.249412234	1.813344516	Up	2.35E-14	8.22E-14	AAEL012741-PA OS=Aedes aegypti GN=AAEL012741 PE=4 SV_1
CCG015482.1	1566	139	540	3.81404413	12.0545782	1.660187728	Up	1.46E-13	4.37E-13	AAEL006290-PA OS=Aedes aegypti GN=AAEL006290 PE=4 SV_1
CCG017308.1	1170	191	540	7.01471736	16.13458928	1.201699973	Up	6.68E-14	2.19E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003916 PE=4 SV_1
CCG015511.1	1863	208	540	4.79747976	10.13283385	1.078689083	Up	3.93E-13	1.05E-12	AAEL006244-PA OS=Aedes aegypti GN=AAEL006244 PE=4 SV_1

CCG014537.1	563	2	541	0.15264559	33.59223399	7.781797988	Up	9.36E-05	0.00015332	Histone H2B OS=Aedes aegypti GN=AAEL015674 PE=3 SV=1
CCG007424.2	957	60	541	2.69402725	19.76220244	2.874907392	Up	0	0	Aldo/keto reductase OS=Aedes aegypti PE=2 SV=1
CCG003751.1	687	76	541	4.75356598	27.52900689	2.533870474	Up	0	0	Putative lipoyltransferase 2, mitochondrial OS=Aedes aegypti GN=AAEL009657 PE=3 SV=1
CCG021078.1	1058	88	541	3.5740422	17.87564058	2.322366369	Up	2.93E-14	1.02E-13	AAEL009459-PA OS=Aedes aegypti GN=AAEL009459 PE=4 SV=1
CCG024281.1	2172	139	541	2.74990475	8.707379251	1.662856915	Up	1.46E-13	4.37E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005947 PE=4 SV=1
CCG004438.1	5810	219	541	1.61968535	3.255151073	1.007010928	Up	4.32E-13	1.14E-12	AAEL000715-PA OS=Aedes aegypti GN=AAEL006715 PE=4 SV=1
CCG014205.1	5256	1002	541	8.19171881	3.598254896	-1.186868806	Down	1.36E-56	6.48E-56	Abc transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ009791 PE=3 SV=1
CCG012548.1	1263	10	542	0.34021959	15.00188916	5.46253415	Up	5.43E-07	1.01E-06	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ019142 PE=4 SV=1
CCG028259.1	1766	93	542	2.26284559	10.72898415	2.245303434	Up	4.04E-14	1.38E-13	AAEL000292-PA OS=Aedes aegypti GN=AAEL000292 PE=4 SV=1
CCG011340.1	630	127	542	8.66215285	30.07521589	1.795777558	Up	1.57E-13	4.67E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003834 PE=4 SV=1
CCG001482.1	813	201	542	10.6235137	23.30551785	1.133410554	Up	0	0	SCO1, mitochondrial OS=Culex quinquefasciatus GN=CpipJ_CPIJ003089 PE=4 SV=1
CCG017255.1	2320	6407	542	118.666849	8.166976729	-3.86097103	Down	0	0	AAEL014553-PA OS=Aedes aegypti GN=AAEL014553 PE=3 SV=1
CCG011881.1	759	32	543	1.81163571	25.00967627	3.787121592	Up	1.77E-13	5.17E-13	AAEL005632-PA OS=Aedes aegypti GN=AAEL005632 PE=4 SV=1
CCG016966.1	561	150	543	11.4892339	33.83662083	1.558302901	Up	0	0	AAEL000780-PA OS=Aedes aegypti GN=AAEL006786 PE=3 SV=1

CCG019048.1	697	161	543	9.92557715	27.23435335	1.456204714	Up	0	0	AAEL009140-PA OS=Aedes aegypti GN=AAEL009140 PE=4 SV=1
CCG024337.1	966	113	544	5.02648034	19.68664862	1.969597083	Up	1.42E-13	4.27E-13	Cuiclicron OS=Culex quinquefasciatus GN=CpipJ_CPIJ005389 PE=4 SV=1
CCG008216.1	1938	114	544	2.52763145	9.81284962	1.956886031	Up	8.48E-14	2.70E-13	Mediator of RNA polymerase II transcription subunit 16 OS=Acromyrmex echinatior GN=G5I_13705 PE=4 SV=1
CCG001966.1	3486	128	544	1.57777568	5.455336364	1.789776045	Up	4.15E-14	1.42E-13	AAEL010924-PA OS=Aedes aegypti GN=AAEL010924 PE=4 SV=1
CCG017567.1	660	170	544	11.0679619	28.81409479	1.380385109	Up	3.53E-13	9.53E-13	AAEL006867-PA OS=Aedes aegypti GN=AAEL006867 PE=4 SV=1
CCG011249.1	35686	1306	544	1.57256272	0.532906534	-1.561163136	Down	9.03E-111	4.86E-110	AAEL006269-PA OS=Aedes aegypti GN=AAEL006269 PE=4 SV=1
CCG004262.1	1083	9	545	0.35708921	17.59211527	5.622500622	Up	1.09E-06	2.01E-06	AAEL009916-PA OS=Aedes aegypti GN=AAEL009916 PE=4 SV=1
CCG017043.1	2592	218	545	3.61396688	7.350409275	1.024241299	Up	0	0	AAEL008807-PA OS=Aedes aegypti GN=AAEL008807 PE=4 SV=1
CCG012843.1	1461	3766	545	110.762506	13.04056183	-3.086391661	Down	0	0	AAEL010951-PA OS=Aedes aegypti GN=AAEL010951 PE=3 SV=1
CCG005005.1	909	55	546	2.59992894	20.99804083	3.013710631	Up	0	0	Nuclear receptor FTZ-F1 protein OS=Aedes aegypti GN=FTZ-F1 PE=2 SV=1
CCG021010.1	1092	58	546	2.28227528	17.47913839	2.93708935	Up	0	0	AAEL003878-PA OS=Aedes aegypti GN=AAEL003878 PE=4 SV=1
CCG016316.1	966	107	546	4.75958758	19.759026	2.053603358	Up	0	0	AAEL008941-PA OS=Aedes aegypti GN=AAEL008941 PE=4 SV=1
CCG026816.1	1287	185	546	6.17669068	14.83078409	1.263688884	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006253 PE=4 SV=1
CCG018190.1	717	941	546	56.3940311	26.62094717	-1.082980568	Down	3.71E-46	1.70E-45	AAEL004728-PA OS=Aedes aegypti GN=AAEL004728 PE=4 SV=1
CCG006509.1	993	15	547	0.64908965	19.25697623	4.890819631	Up	1.48E-08	2.93E-08	AAEL000698-PA OS=Aedes aegypti GN=AAEL000698 PE=4 SV=1

CCG005079.1	1206	89	547	3.17106665	15.85586849	2.321976796	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_15285 PE=4 SV=1
CCG025104.1	1281	113	547	3.79046059	14.92753895	1.977531264	Up	1.42E-13	4.27E-13	AAEL002129-PA OS=Aedes aegypti GN=AAEL002129 PE=4 SV=1
CCG025833.1	6669	5142	547	33.1309605	2.867323046	-3.530403668	Down	0	0	AAEL002228-PA OS=Aedes aegypti GN=AAEL002228 PE=4 SV=1
CCG006088.1	1563	0	548	0.001	12.2566447	13.58127647	Up	4.44E-16	1.66E-15	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG023770.1	2146	12	548	0.24027811	8.926903854	5.215382786	Up	1.30E-07	2.49E-07	AAEL001635-PA OS=Aedes aegypti GN=AAEL001635 PE=4 SV=1
CCG010101.1	1503	38	548	1.08639382	12.74593192	3.552417773	Up	4.26E-14	1.45E-13	AAEL012851-PA OS=Aedes aegypti GN=AAEL012851 PE=4 SV=1
CCG001663.1	414	99	548	10.2753713	46.27327457	2.170988667	Up	0	0	-
CCG020685.1	4077	106	548	1.11719202	4.698831413	2.072424832	Up	0	0	AAEL008592-PA OS=Aedes aegypti GN=AAEL008592 PE=3 SV=1
CCG010181.1	1977	145	548	3.15154857	9.690002869	1.620436197	Up	0	0	AAEL006459-PA OS=Aedes aegypti GN=AAEL006459 PE=4 SV=1
CCG010204.2	1965	148	548	3.23639731	9.749178459	1.590891921	Up	0	0	AAEL001729-PA OS=Aedes aegypti GN=AAEL001729 PE=4 SV=1
CCG016294.1	1016	189	548	7.99338567	18.8554485	1.238102863	Up	2.34E-13	6.64E-13	AAEL001105-PA OS=Aedes aegypti GN=AAEL001105 PE=4 SV=1
CCG026613.1	2619	1237	548	20.2953653	7.314675705	-1.472284498	Down	2.23E-96	1.17E-95	AAEL001320-PA OS=Aedes aegypti GN=CYP9M4 PE=3 SV=1
CCG001672.1	984	22	549	0.96070545	19.50416052	4.343543924	Up	8.38E-11	1.80E-10	AAEL012018-PA OS=Aedes aegypti GN=AAEL012018 PE=4 SV=1
CCG011298.1	3741	47	549	0.53984965	5.13020421	3.248386691	Up	0	0	AAEL010940-PA (Fragment) OS=Aedes aegypti GN=AAEL010940 PE=4 SV=1
CCG009266.1	498	163	549	14.064391	38.53834126	1.454247389	Up	1.80E-13	5.24E-13	AAEL003180-PA OS=Aedes aegypti GN=AAEL003180 PE=4 SV=1
CCG007239.1	1986	180	549	3.89453788	9.663692824	1.311122446	Up	3.32E-13	9.05E-13	AAEL006362-PA OS=Aedes aegypti GN=AAEL006362 PE=3 SV=1
CCG022399.1	5607	23735	549	181.895247	3.422881032	-5.731752886	Down	0	0	AAEL005416-PA OS=Aedes aegypti GN=AAEL005416 PE=4 SV=1

CCG006983.1	1088	17	550	0.6714021	17.671923	4.718138171	Up	3.39E-09	6.89E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002513 PE=4 SV=1
CCG009017.1	836	82	550	4.21473473	22.9988663	2.448049008	Up	0	0	AAEL014657-PA OS=Aedes aegypti GN=AAEL014657 PE=4 SV=1
CCG020522.1	999	92	550	3.95717276	19.24629852	2.282039056	Up	1.44E-14	5.11E-14	AAEL008561-PA OS=Aedes aegypti GN=AAEL008561 PE=4 SV=1
CCG026454.1	966	116	550	5.15992672	19.90378077	1.947620017	Up	1.16E-13	3.59E-13	AAEL003202-PA OS=Aedes aegypti GN=AAEL003202 PE=4 SV=1
CCG006248.1	910	204	550	9.63277567	21.12862882	1.13317567	Up	6.59E-14	2.17E-13	cGMP-phosphodiesterase delta subunit OS=Glossina morsitans morsitans PE=2 SV=1
CCG003024.1	792	210	550	11.3934902	24.27658109	1.091355495	Up	0	0	AAEL010803-PA (Fragment) OS=Aedes aegypti GN=AAEL010803 PE=3 SV=1
CCG014709.2	1309	20	551	0.65652765	14.71505768	4.486293618	Up	3.69E-10	7.71E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010882 PE=4 SV=1
CCG016515.1	1016	67	551	2.83363407	18.95867175	2.742132522	Up	0	0	AAEL006177-PA OS=Aedes aegypti GN=AAEL006177 PE=4 SV=1
CCG025747.1	591	107	551	7.77963046	32.59223435	2.066754726	Up	0	0	AAEL012990-PA OS=Aedes aegypti GN=AAEL012990 PE=4 SV=1
CCG000822.1	615	134	551	9.36251128	31.32034228	1.742132522	Up	0	0	AAEL006654-PA OS=Aedes aegypti GN=AAEL006654 PE=4 SV=1
CCG009433.1	1312	161	551	5.27296286	14.68141044	1.477304834	Up	0	0	AAEL003084-PA (Fragment) OS=Aedes aegypti GN=AAEL801117_b PE=2 SV=1
CCG002892.1	777	189	551	10.4520976	24.79023231	1.245979288	Up	2.34E-13	6.65E-13	AAEL011810-PA OS=Aedes aegypti GN=AAEL011810 PE=4 SV=1
CCG003412.1	1770	4274	551	103.758557	10.88249181	-3.25314948	Down	0	0	AAEL004366-PA OS=Aedes aegypti GN=AAEL004366 PE=4 SV=1
CCG010907.1	467	6	552	0.55207368	41.32113229	6.22587516	Up	8.38E-06	1.47E-05	Histone H2A OS=Gasterosteus aculeatus PE=3 SV=1
CCG015016.1	940	13	552	0.59426229	20.52869019	5.110397942	Up	6.34E-08	1.23E-07	AAEL010667-PA OS=Aedes aegypti GN=AAEL010667 PE=3 SV=1

CCG019027.1	966	47	552	2.09065997	19.97615816	3.256248809	Up	0	0	AAEL008833-PA OS=Aedes aegypti GN=AAEL008833 PE=4 SV=1
CCG004022.2	655	78	552	5.11700656	29.46102104	2.525435442	Up	0	0	AAEL007790-PA OS=Aedes aegypti GN=AAEL007790 PE=4 SV=1
CCG000134.1	1814	118	552	2.79516465	10.63779977	1.928194611	Up	0	0	AAEL002426-PA OS=Aedes aegypti GN=AAEL002426 PE=4 SV=1
CCG016669.1	862	166	552	8.27491409	22.38627468	1.435798229	Up	1.83E-13	5.32E-13	AAEL000410-PA OS=Aedes aegypti GN=AAEL000410 PE=3 SV=1
CCG025984.1	1569	194	552	5.31302009	12.29889661	1.210924818	Up	5.97E-13	1.53E-12	AAEL002013-PA OS=Aedes aegypti GN=AAEL002013 PE=4 SV=1
CCG018349.1	2766	973	552	15.1155285	6.976489074	-1.115458334	Down	6.20E-50	2.89E-49	AAEL008629-PA OS=Aedes aegypti GN=AAEL008629 PE=3 SV=1
CCG010649.1	464	1	553	0.09260719	41.66363589	8.813448874	Up	0.000587	0.00091581	Histone H2A OS=Gasterosteus aculeatus PE=3 SV=1
CCG018544.1	3411	107	553	1.3479219	5.667524789	2.071981888	Up	0	0	AAEL010928-PA (Fragment) OS=Aedes aegypti GN=AAEL010928 PE=4 SV=1
CCG001269.1	8017	1492	553	7.99686217	2.411366728	-1.729582946	Down	3.33E-146	1.89E-145	AAEL009173-PA OS=Aedes aegypti GN=AAEL009173 PE=4 SV=1
CCG001804.1	1371	4	554	0.12536757	14.12610163	6.81605537	Up	2.95E-05	5.03E-05	AAEL013582-PA OS=Aedes aegypti GN=AAEL013582 PE=4 SV=1
CCG021990.1	1653	4	554	0.10398	11.71620407	6.81605537	Up	2.95E-05	5.03E-05	AAEL006333-PA OS=Aedes aegypti GN=AAEL006333 PE=3 SV=1
CCG014241.1	1108	86	554	3.33519601	17.47913839	2.389790615	Up	0	0	AAEL006839-PA OS=Aedes aegypti GN=AAEL006839 PE=4 SV=1
CCG024347.1	2386	4100	554	73.8373478	8.116884045	-3.185352825	Down	0	0	AAEL005249-PA OS=Aedes aegypti GN=AAEL005249 PE=3 SV=1
CCG005416.1	1293	21	555	0.69788432	15.00529281	4.426339742	Up	1.76E-10	3.72E-10	AAEL003773-PA OS=Aedes aegypti GN=AAEL003773 PE=4 SV=1
CCG011513.1	1710	30	555	0.75385499	11.34610737	3.911766569	Up	3.71E-13	9.95E-13	AAEL007045-PA OS=Aedes aegypti GN=AAEL007045 PE=4 SV=1
CCG023392.1	1475	47	555	1.3692051	13.15379228	3.264068313	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_05553 PE=4 SV=1

CCG016435.2	1326	143	555	4.63399099	14.63185793	1.658785828	Up	0	0	AAEL003650-PA OS=Aedes aegypti GN=AAEL003650 PE=4 SV=1
CCG015756.1	2195	2369	555	46.3759915	8.839108706	-2.391405318	Down	0	0	AAEL000992-PA OS=Aedes aegypti GN=CYP6AG6 PE=3 SV=1
CCG027682.1	579	0	556	0.001	33.56960602	15.03486798	Up	4.44E-16	1.65E-15	AAEL000655-PA OS=Aedes aegypti GN=AAEL008835 PE=4 SV=1
CCG011766.1	5769	135	556	1.00553201	3.369180427	1.744438679	Up	1.90E-13	5.48E-13	AAEL012505-PA OS=Aedes aegypti GN=AAEL012505 PE=4 SV=1
CCG003698.1	1862	185	556	4.26928083	10.43866911	1.289872816	Up	0	0	AAEL005057-PA OS=Aedes aegypti GN=AAEL005057 PE=4 SV=1
CCG017629.1	675	3145	556	200.207134	28.79526205	-2.797590025	Down	0	0	AAEL004987-PA OS=Aedes aegypti GN=AAEL004987 PE=4 SV=1
CCG001479.1	1344	42	557	1.34280421	14.48791679	3.431529298	Up	7.66E-14	2.48E-13	AAEL010138-PA OS=Aedes aegypti GN=AAEL010138 PE=4 SV=1
CCG011148.1	3114	60	557	0.82793323	6.252973719	2.916956126	Up	0	0	AAEL005888-PA OS=Aedes aegypti GN=AAEL005888 PE=4 SV=1
CCG002514.1	1809	73	557	1.7339915	10.76382541	2.634022162	Up	0	0	AAEL005478-PA (Fragment) OS=Aedes aegypti GN=AAEL005478 PE=4 SV=1
CCG025433.1	2637	78	557	1.27100466	7.384057703	2.538444502	Up	0	0	AAEL005438-PA OS=Aedes aegypti GN=AAEL005438 PE=4 SV=1
CCG024753.1	981	1	558	0.04380197	19.8845244	8.826434516	Up	0.000587	0.00091802	AAEL000684-PA OS=Aedes aegypti GN=AAEL006884 PE=4 SV=1
CCG025254.1	1651	43	558	1.11913906	11.81509294	3.400169761	Up	8.46E-14	2.70E-13	AAEL007471-PA OS=Aedes aegypti GN=AAEL007471 PE=4 SV=1
CCG027946.1	1274	80	558	2.69825649	15.31139595	2.504506421	Up	0	0	AAEL009611-PA OS=Aedes aegypti GN=AAEL009611 PE=4 SV=1
CCG020967.1	2214	131	558	2.54247301	8.810622601	1.793011514	Up	1.26E-13	3.87E-13	AAEL009383-PA OS=Aedes aegypti GN=AAEL009383 PE=4 SV=1
CCG017047.1	660	147	558	9.5705318	29.555634	1.626762171	Up	7.02E-14	2.28E-13	AAEL008800-PA OS=Aedes aegypti GN=AAEL008800 PE=4 SV=1
CCG011966.1	2853	958	558	14.4286736	6.837265489	-1.07744733	Down	1.39E-46	6.42E-46	AAEL003076-PA OS=Aedes aegypti GN=AAEL003076 PE=4 SV=1
CCG008834.1	621	92	559	6.36588661	31.46807845	2.305455721	Up	1.44E-14	5.12E-14	AAEL013778-PA OS=Aedes aegypti GN=AAEL013778 PE=4 SV=1

CCG012450.2	1841	119	559	2.77751136	10.61470761	1.934199913	Up	0	0	AAEL011904-PA OS=Aedes aegypti GN=AAEL011904 PE=4 SV=1
CCG005859.1	495	125	559	10.8509431	39.47813478	1.863233392	Up	2.35E-14	8.22E-14	AAEL009276-PA OS=Aedes aegypti GN=AAEL009276 PE=4 SV=1
CCG017162.1	1356	136	559	4.3096489	14.41126601	1.741554835	Up	1.70E-13	4.99E-13	AAEL000477-PA OS=Aedes aegypti GN=AAEL000477 PE=4 SV=1
CCG004549.1	657	161	559	10.5298741	29.74380018	1.498100799	Up	0	0	AAEL011720-PA OS=Aedes aegypti GN=AAEL011720 PE=4 SV=1
CCG003191.1	2058	23	560	0.48022541	9.512456265	4.308034265	Up	4.01E-11	8.68E-11	AAEL000952-PA OS=Aedes aegypti GN=AAEL000952 PE=4 SV=1
CCG009836.1	2963	48	560	0.696101	6.607031722	3.24663372	Up	5.35E-14	1.79E-13	Zinc finger protein 557 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014708 PE=4 SV=1
CCG026677.1	2189	57	560	1.11890127	8.943186383	2.998706207	Up	0	0	AAEL003949-PA OS=Aedes aegypti GN=AAEL003949 PE=4 SV=1
CCG021613.1	1935	62	560	1.37680803	10.11712403	2.87739991	Up	0	0	AAEL002366-PA OS=Aedes aegypti GN=AAEL002366 PE=4 SV=1
CCG013809.1	1023	140	560	5.88051109	19.13649559	1.702313204	Up	0	0	Putative arsenite-translocating ATPase OS=Glossina morsitans morsitans PE=2 SV=1
CCG009389.1	2649	178	560	2.88735854	7.390198185	1.35586279	Up	1.15E-13	3.58E-13	AAEL008583-PA OS=Aedes aegypti GN=AAEL008583 PE=4 SV=1
CCG013408.2	1446	215	560	6.38899927	13.5384751	1.083403371	Up	0	0	AAEL004553-PA OS=Aedes aegypti GN=AAEL004553 PE=4 SV=1
CCG019110.1	794	3	561	0.16235416	24.69973963	7.249207664	Up	5.28E-05	8.85E-05	-
CCG007408.1	1441	62	561	1.84880191	13.6097108	2.879973854	Up	0	0	AAEL004053-PA OS=Aedes aegypti GN=AAEL004053 PE=4 SV=1
CCG020738.1	846	1	562	0.05079165	23.22287417	8.836739524	Up	0.000587	0.00091808	AAEL010318-PA OS=Aedes aegypti GN=AAEL010318 PE=2 SV=1
CCG002106.1	2081	51	562	1.05307855	9.440918571	3.164314182	Up	0	0	AAEL003176-PA OS=Aedes aegypti GN=AAEL003176 PE=4 SV=1
CCG019114.1	351	57	562	6.97799109	55.97308133	3.00384951	Up	0	0	-
CCG014757.1	1881	21	563	0.4797259	10.46332261	4.446986893	Up	1.76E-10	3.73E-10	AAEL002647-PA OS=Aedes aegypti GN=AAEL002647 PE=4 SV=1

CCG004224.1	913	33	563	1.55312294	21.55696585	3.794910197	Up	1.31E-13	4.02E-13	AAEL011866-PA OS=Aedes aegypti GN=AAEL011866 PE=4 SV=1
CCG019242.1	570	45	564	3.39234747	34.59029491	3.35001146	Up	0	0	AAEL001399-PA OS=Aedes aegypti GN=AAEL001399 PE=4 SV=1
CCG007619.1	3681	95	564	1.10897169	5.356280386	2.272008948	Up	0	0	AAEL010370-PA OS=Aedes aegypti GN=AAEL010370 PE=4 SV=1
CCG001919.1	2024	96	564	2.03809018	9.741337994	2.256902055	Up	0	0	AAEL001585-PA (Fragment) OS=Aedes aegypti GN=AAEL001585 PE=4 SV=1
CCG012044.1	3528	100	564	1.217963	5.588568055	2.198008366	Up	0	0	AAEL003089-PA OS=Aedes aegypti GN=AAEL003089 PE=4 SV=1
CCG019901.1	1611	120	564	3.20072511	12.23865183	1.934973961	Up	0	0	AAEL002384-PA OS=Aedes aegypti GN=AAEL002384 PE=4 SV=1
CCG003022.1	3426	153	564	1.91896363	5.754952744	1.584476714	Up	0	0	AAEL010796-PA OS=Aedes aegypti GN=AAEL010796 PE=4 SV=1
CCG021821.1	2923	187	564	2.74900457	6.745285015	1.294970096	Up	3.31E-13	9.03E-13	AAEL005892-PA OS=Aedes aegypti GN=AAEL005892 PE=4 SV=1
CCG023499.1	609	38	565	2.68119855	32.43255563	3.596492748	Up	4.26E-14	1.45E-13	AAEL005216-PA OS=Aedes aegypti GN=AAEL005216 PE=3 SV=1
CCG003018.1	1871	62	565	1.42390355	10.55661485	2.890223951	Up	0	0	AAEL010005-PA OS=Aedes aegypti GN=AAEL010005 PE=4 SV=1
CCG006865.4	1107	2735	565	106.162804	17.84230025	-2.572904857	Down	0	0	AAEL007874-PA OS=Aedes aegypti GN=AAEL007874 PE=4 SV=1
CCG009141.1	6516	37	566	0.24399634	3.036584508	3.637518081	Up	5.17E-14	1.74E-13	DNA polymerase OS=Aedes aegypti GN=AAEL002800 PE=3 SV=1
CCG020941.1	568	42	566	3.17733953	34.83518425	3.454654024	Up	7.66E-14	2.47E-13	AAEL008910-PA OS=Aedes aegypti GN=AAEL008910 PE=4 SV=1
CCG004504.1	1303	56	566	1.84674224	15.185253	3.039616525	Up	0	0	Integrator complex subunit 12 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008030 PE=4 SV=1
CCG008008.1	447	95	566	9.13227022	44.26484262	2.277115838	Up	0	0	AAEL003412-PA OS=Aedes aegypti GN=AAEL003412 PE=4 SV=1
CCG012885.1	1105	133	566	5.17192281	17.90623046	1.791689011	Up	9.90E-14	3.11E-13	AAEL009501-PA OS=Aedes aegypti GN=AAEL009501 PE=3 SV=1

CCG016108.1	780	2162	566	119.10329	25.36715981	-2.231179361	Down	2.23E-296	1.40E-295	AAEL014366-PA OS=Aedes aegypti GN=AAEL014366 PE=4 SV=1
CCG007300.1	271	20	567	3.17119813	73.14148683	4.527590034	Up	3.69E-10	7.71E-10	AAEL005428-PA OS=Aedes aegypti GN=AAEL005428 PE=4 SV=1
CCG021720.1	594	82	567	5.93184889	33.36926419	2.491966124	Up	0	0	-
CCG011773.1	934	133	567	6.1188166	21.22199457	1.794235693	Up	9.90E-14	3.11E-13	AAEL002374-PA OS=Aedes aegypti GN=AAEL002374 PE=4 SV=1
CCG026146.1	2210	158	567	3.07204437	8.968933452	1.545737381	Up	0	0	AAEL009512-PA OS=Aedes aegypti GN=AAEL009512 PE=4 SV=1
CCG002236.1	4092	161	567	1.69064694	4.843925447	1.518601251	Up	0	0	Num1-associated kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ004075 PE=4 SV=1
CCG021752.1	993	175	567	7.57271254	19.96107042	1.398307017	Up	3.55E-13	9.58E-13	AAEL003230-PA OS=Aedes aegypti GN=AAEL003230 PE=4 SV=1
CCG018477.1	2001	1133	567	24.3301896	9.905718606	-1.296414017	Down	4.70E-73	2.35E-72	AAEL007153-PA OS=Aedes aegypti GN=AAEL007153 PE=4 SV=1
CCG028062.1	1803	12	568	0.28598825	11.01292358	5.267097823	Up	1.30E-07	2.49E-07	AAEL000904-PA (Fragment) OS=Aedes aegypti GN=AAEL000904 PE=4 SV=1
CCG014882.1	1583	89	568	2.41586	12.54346254	2.376326892	Up	0	0	AAEL011765-PA OS=Aedes aegypti GN=AAEL011765 PE=4 SV=1
CCG023184.1	1031	102	568	4.25112796	19.25926402	2.179634981	Up	0	0	AAEL013998-PA OS=Aedes aegypti GN=AAEL013998 PE=4 SV=1
CCG008695.1	1083	116	568	4.60248312	18.33453482	1.994079328	Up	1.16E-13	3.59E-13	AAEL009916-PA OS=Aedes aegypti GN=AAEL009916 PE=4 SV=1
CCG003353.1	1238	193	568	6.69883585	16.03901551	1.259603286	Up	6.08E-14	2.02E-13	AAEL005573-PA OS=Aedes aegypti GN=AAEL005573 PE=4 SV=1
CCG001079.1	603	1050	568	74.822921	32.9291894	-1.184113289	Down	5.26E-59	2.52E-58	AAEL008054-PA OS=Aedes aegypti GN=AAEL008054 PE=4 SV=1
CCG019729.1	847	29	569	1.47121878	23.48436775	3.996617051	Up	6.66E-13	1.69E-12	AAEL010667-PA OS=Aedes aegypti GN=AAEL010667 PE=3 SV=1
CCG004879.1	484	144	569	12.7843839	41.09764356	1.684673045	Up	7.53E-14	2.43E-13	AAEL005841-PA OS=Aedes aegypti GN=AAEL005841 PE=4 SV=1
CCG025115.1	1515	197	569	5.58748364	13.12954421	1.232546227	Up	0	0	AAEL003471-PA OS=Aedes aegypti GN=AAEL003471 PE=4 SV=1

CCG023825.1	1019	989	569	41.7046786	19.52037241	-1.095228665	Down	2.74E-49	1.28E-48	AAEL001403-PA OS=Aedes aegypti GN=AAEL001403 PE=4 SV=1
CCG009450.1	1510	1252	569	35.6278859	13.17301953	-1.435420801	Down	6.38E-94	3.34E-93	AAEL007410-PA OS=Aedes aegypti GN=AAEL007410 PE=4 SV=1
CCG003391.1	975	101	570	4.45122379	20.43714642	2.19891983	Up	0	0	AAEL014649-PA OS=Aedes aegypti GN=AAEL014649 PE=4 SV=1
CCG003667.2	1146	120	570	4.49944865	17.38762457	1.950240717	Up	0	0	AAEL000204-PA OS=Aedes aegypti GN=AAEL000204 PE=4 SV=1
CCG022653.1	2313	167	570	3.10244085	8.614880138	1.47342702	Up	1.62E-13	4.78E-13	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1
CCG018862.1	1743	186	570	4.58541058	11.4321387	1.317972502	Up	6.43E-13	1.64E-12	AAEL011872-PA OS=Aedes aegypti GN=AAEL011872 PE=4 SV=1
CCG019412.1	732	3231	570	189.665591	27.22160896	-2.800633722	Down	0	0	-
CCG014542.1	2733	6	571	0.09433531	7.303759984	6.274697638	Up	8.38E-06	1.48E-05	AAEL009787-PA OS=Aedes aegypti GN=AAEL009787 PE=4 SV=1
CCG004560.1	604	23	571	1.63626473	33.0483047	4.336098183	Up	4.01E-11	8.68E-11	Uncharacterized protein (Fragment) OS=Monodelphis domestica PE=4 SV=1
CCG000835.1	4443	86	571	0.83173468	4.492724744	2.433395384	Up	0	0	Chromatin regulatory protein sir2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006726 PE=4 SV=1
CCG020808.1	1530	110	571	3.08932732	13.04652029	2.078300426	Up	0	0	AAEL001190-PA OS=Aedes aegypti GN=AAEL001190 PE=4 SV=1
CCG019514.1	1785	115	571	2.76835825	11.18273167	2.014170088	Up	6.13E-14	2.03E-13	AAEL013355-PA OS=Aedes aegypti GN=AAEL013355 PE=4 SV=1
CCG015863.1	1052	127	571	5.18741093	18.97450194	1.870975452	Up	1.57E-13	4.67E-13	AAEL007647-PA OS=Aedes aegypti GN=AAEL007647 PE=4 SV=1
CCG023088.1	3078	204	571	2.84789664	6.485112423	1.187234797	Up	6.59E-14	2.17E-13	AAEL014815-PA (Fragment) OS=Aedes aegypti GN=AAEL014815 PE=4 SV=1
CCG018522.1	1862	206	571	4.7539019	10.72028788	1.173159612	Up	1.10E-13	3.43E-13	AAEL004538-PA OS=Aedes aegypti GN=AAEL004538 PE=3 SV=1
CCG003758.1	2385	1142	571	20.575026	8.369465844	-1.297686796	Down	9.92E-74	4.97E-73	AAEL003413-PA OS=Aedes aegypti GN=AAEL003413 PE=4 SV=1

CCG025893.1	915	0	572	0.001	21.8536987	14.41558985	Up	4.44E-16	1.64E-15	AAEL010638-PA OS=Aedes aegypti GN=AAEL010638 PE=3 SV=1
CCG019129.1	753	6	572	0.34238832	26.55529125	6.27722204	Up	8.38E-06	1.48E-05	AAEL009962-PA OS=Aedes aegypti GN=AAEL009962 PE=4 SV=1
CCG009602.1	1174	37	572	1.35424206	17.03248238	3.652731175	Up	5.17E-14	1.74E-13	AAEL005592-PA OS=Aedes aegypti GN=AAEL005592 PE=4 SV=1
CCG017681.1	2187	59	572	1.15922009	9.143179842	2.979541491	Up	0	0	Condensin complex subunit 2 OS=Aedes aegypti GN=AAEL005962 PE=3 SV=1
CCG018433.1	1530	61	572	1.71317243	13.06936883	2.931447203	Up	1.75E-14	6.17E-14	Zinc finger protein 8 OS=Culex quinquefasciatus GN=CpipJ_CPIJ012305 PE=4 SV=1
CCG004049.1	1254	80	572	2.74129088	15.94588063	2.540256446	Up	0	0	AAEL009429-PA (Fragment) OS=Aedes aegypti GN=AAEL009429 PE=4 SV=1
CCG000805.1	945	131	572	5.95665104	21.15993049	1.828761539	Up	1.26E-13	3.87E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003895 PE=4 SV=1
CCG003317.1	2070	133	572	2.76085734	9.659968268	1.806902105	Up	9.90E-14	3.11E-13	AAEL002563-PA OS=Aedes aegypti GN=AAEL002563 PE=4 SV=1
CCG025489.1	531	182	572	14.7278563	37.65750342	1.3543899	Up	0	0	AAEL000825-PA OS=Aedes aegypti GN=AAEL000825 PE=4 SV=1
CCG012486.1	1131	193	572	7.33258955	17.68004802	1.269727503	Up	6.08E-14	2.02E-13	AAEL008408-PA OS=Aedes aegypti GN=AAEL008408 PE=4 SV=1
CCG027211.1	4743	2539	572	23.0023521	4.21592543	-2.447860139	Down	0	0	AAEL005416-PA OS=Aedes aegypti GN=AAEL005416 PE=4 SV=1
CCG021973.1	5703	8774	572	66.1084432	3.506248345	-4.23683445	Down	0	0	Uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ019410 PE=4 SV=1
CCG018500.1	742	2	573	0.11582139	26.99608166	7.864704533	Up	9.36E-05	0.00015376	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG000059.1	2352	72	573	1.31540004	8.516620999	2.694779531	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_20768 PE=4 SV=1

CCG000754.1	3159	116	573	1.57786933	6.340959984	2.006723537	Up	1.16E-13	3.59E-13	AAEL013736-PA (Fragment) OS=Aedes aegypti GN=AAEL013736 PE=4 SV=1
CCG010846.1	1305	122	573	4.01709396	15.34949624	1.933967195	Up	8.04E-14	2.58E-13	Polyadenylate-binding protein-like protein OS=Aedes aegypti PE=2 SV=1
CCG006824.1	2672	123	573	1.97802296	7.496666389	1.922190027	Up	0	0	AAEL002998-PA OS=Aedes aegypti GN=AAEL002998 PE=4 SV=1
CCG014825.1	1932	193	573	4.29252525	10.36806035	1.272247495	Up	6.08E-14	2.02E-13	AAEL010246-PA OS=Aedes aegypti GN=AAEL010246 PE=4 SV=1
CCG020642.1	1043	33	574	1.35954098	19.23878319	3.822826011	Up	1.31E-13	4.01E-13	AAEL011383-PA OS=Aedes aegypti GN=AAEL011383 PE=4 SV=1
CCG003785.1	1715	50	574	1.25276194	11.70032121	3.223363941	Up	0	0	AAEL010013-PA OS=Aedes aegypti GN=AAEL010013 PE=4 SV=1
CCG006498.2	3589	72	574	0.86202867	5.590986589	2.697295129	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006312 PE=4 SV=1
CCG017467.1	1544	111	574	3.08914543	12.99614693	2.072804264	Up	0	0	AAEL009476-PA OS=Aedes aegypti GN=AAEL009476 PE=4 SV=1
CCG013995.1	1428	123	574	3.70117462	14.05185635	1.924705625	Up	0	0	AAEL003762-PA OS=Aedes aegypti GN=AAEL003762 PE=4 SV=1
CCG027340.1	1710	206	574	5.17647095	11.73453267	1.180719603	Up	1.10E-13	3.43E-13	AAEL002837-PA OS=Aedes aegypti GN=AAEL002837 PE=4 SV=1
CCG009123.1	1149	70	575	2.61782543	17.49435087	2.740448333	Up	0	0	AAEL003124-PA OS=Aedes aegypti GN=AAEL003124 PE=4 SV=1
CCG020229.1	396	92	575	9.98286764	50.7601241	2.346169394	Up	1.44E-14	5.11E-14	AAEL000435-PA OS=Aedes aegypti GN=AAEL000435 PE=4 SV=1
CCG014500.1	1020	139	575	5.85567952	19.70687171	1.750790277	Up	1.46E-13	4.37E-13	AAEL006342-PA OS=Aedes aegypti GN=AAEL006342 PE=4 SV=1
CCG027655.1	1961	146	575	3.19917453	10.25038712	1.679906791	Up	9.33E-15	3.34E-14	AAEL013208-PA OS=Aedes aegypti GN=AAEL013208 PE=4 SV=1
CCG007432.1	2981	1481	575	21.3479292	6.743042316	-1.662624576	Down	2.95E-137	1.66E-136	AAEL004090-PA (Fragment) OS=Aedes aegypti GN=AAEL004090 PE=4 SV=1

CCG011095.1	1690	49	576	1.24586804	11.91477362	3.257528361	Up	0	0	AAEL005202-PA OS=Reedes aegypti GN=AAEL005202 PE=4 SV=1
CCG020661.1	1793	70	576	1.67756911	11.23032204	2.742955188	Up	0	0	AGAP009012-PA OS=Anopheles gambiae GN=AGAP009012 PE=4 SV=4
CCG019538.3	2271	83	576	1.57044825	8.866564254	2.497198774	Up	1.32E-13	4.02E-13	AAEL008721-PA OS=Reedes aegypti GN=AAEL008721 PE=4 SV=1
CCG013626.1	900	96	576	4.58343836	22.37329713	2.287275705	Up	0	0	AAEL005851-PA OS=Reedes aegypti GN=AAEL005851 PE=4 SV=1
CCG017239.2	1617	98	576	2.60422634	12.45267002	2.257528361	Up	0	0	AAEL001212-PA OS=Reedes aegypti GN=AAEL001212 PE=4 SV=1
CCG007687.1	2962	147	576	2.13252903	6.798098387	1.67256586	Up	7.02E-14	2.29E-13	AAEL011415-PA OS=Reedes aegypti GN=AAEL011415 PE=4 SV=1
CCG022319.1	1554	69	577	1.90792258	12.98000367	2.766216256	Up	0	0	AAEL003899-PA OS=Reedes aegypti GN=AAEL003899 PE=4 SV=1
CCG023774.1	1386	119	577	3.68932065	14.55333744	1.979922949	Up	0	0	AAEL001638-PA OS=Reedes aegypti GN=AAEL001638 PE=4 SV=1
CCG015493.1	477	6680	577	601.756451	42.28705597	-3.830891675	Down	0	0	AAEL005768-PA OS=Reedes aegypti GN=AAEL005768 PE=4 SV=1
CCG005908.1	570	44	578	3.31696197	35.44891925	3.417807268	Up	3.91E-14	1.34E-13	AAEL010085-PA OS=Reedes aegypti GN=AAEL010085 PE=4 SV=1
CCG017276.1	965	173	578	7.70338247	20.93873987	1.442610659	Up	3.60E-13	9.69E-13	AAEL010964-PA OS=Reedes aegypti GN=AAEL010964 PE=4 SV=1
CCG011302.1	1584	228	578	6.18503756	12.75623988	1.044348872	Up	1.05E-13	3.28E-13	AAEL010938-PA OS=Reedes aegypti GN=AAEL010938 PE=4 SV=1
CCG018229.1	773	9772	578	543.208598	26.1395653	-4.377199262	Down	0	0	Myosin light chain 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017123 PE=4 SV=1
CCG020636.1	1203	65	579	2.32172298	16.82530528	2.857364929	Up	0	0	AAEL001407-PA OS=Reedes aegypti GN=AAEL001407 PE=4 SV=1
CCG026191.1	1278	75	579	2.52169804	15.83790473	2.650914051	Up	0	0	AAEL014105-PA OS=Reedes aegypti GN=AAEL014105 PE=4 SV=1
CCG004895.1	2895	128	579	1.89987082	6.991655354	1.879732742	Up	4.15E-14	1.42E-13	AAEL011693-PA OS=Reedes aegypti GN=AAEL011693 PE=3 SV=1

CCG010339.1	2838	159	579	2.40739528	7.132079722	1.566849787	Up	5.53E-14	1.84E-13	AAEL003865-PA OS=Aedes aegypti GN=AAEL003865 PE=4 SV=1
CCG014586.1	2902	1412	579	20.9073967	6.974790576	-1.583791631	Down	3.95E-122	2.17E-121	AAEL009937-PA OS=Aedes aegypti GN=AAEL009937 PE=4 SV=1
CCG009828.1	1398	39	580	1.1987265	14.50343385	3.596820075	Up	7.33E-14	2.38E-13	AAEL014400-PA OS=Aedes aegypti GN=AAEL014400 PE=4 SV=1
CCG007386.1	1602	53	580	1.42159546	12.65655464	3.154301839	Up	0	0	AAEL007654-PA OS=Aedes aegypti GN=AAEL007654 PE=4 SV=1
CCG004213.1	872	56	580	2.75952424	23.25206483	3.074867372	Up	0	0	AAEL000110-PA OS=Aedes aegypti GN=AAEL000110 PE=4 SV=1
CCG017094.1	1215	183	580	6.47198472	16.68790167	1.366522456	Up	3.05E-13	8.41E-13	AAEL003346-PA OS=Aedes aegypti GN=AAEL003346 PE=4 SV=1
CCG009370.1	1860	209	580	4.82831964	10.90096803	1.174863162	Up	0	0	AAEL007248-PA OS=Aedes aegypti GN=AAEL007248 PE=4 SV=1
CCG005175.1	546	227	580	17.8647065	37.1351658	1.055673807	Up	0	0	AAEL008710-PA OS=Aedes aegypti GN=AAEL008710 PE=4 SV=1
CCG023424.1	4922	1171	580	10.2229905	4.119423106	-1.311303067	Down	9.09E-77	4.58E-76	-
CCG016266.1	1248	1609	580	55.3992812	16.24663504	-1.769726317	Down	1.18E-162	6.82E-162	AAEL007128-PA OS=Aedes aegypti GN=AAEL007128 PE=4 SV=1
CCG027473.1	4486	61	581	0.58429644	4.527587785	2.95397022	Up	1.75E-14	6.18E-14	AAEL008475-PA OS=Aedes aegypti GN=AAEL008475 PE=4 SV=1
CCG016033.1	1319	117	581	3.81156857	15.3986041	2.014342838	Up	9.44E-14	2.97E-13	AAEL002069-PA OS=Aedes aegypti GN=AAEL002069 PE=4 SV=1
CCG023373.1	1800	153	581	3.65242744	11.28375489	1.627319715	Up	0	0	AAEL010598-PA OS=Aedes aegypti GN=AAEL010598 PE=4 SV=1
CCG006149.1	2013	83	582	1.77172776	10.10716199	2.512149115	Up	1.32E-13	4.01E-13	AAEL002013-PA OS=Aedes aegypti GN=AAEL002013 PE=4 SV=1
CCG015581.1	1720	101	582	2.52322279	11.82890528	2.228977064	Up	0	0	AAEL005677-PA OS=Aedes aegypti GN=AAEL005677 PE=4 SV=1
CCG003531.1	1114	152	582	5.86301585	18.26365986	1.639261033	Up	8.39E-14	2.69E-13	SNARE_virga_protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007069 PE=4 SV=1
CCG016959.1	1524	1067	582	30.0844533	13.35020806	-1.172155914	Down	5.42E-59	2.60E-58	AAEL004496-PA OS=Aedes aegypti GN=AAEL004496 PE=4 SV=1

CCG011262.1	4161	1512	582	15.6140925	4.889621985	-1.675053877	Down	1.40E-141	7.87E-141	AAEL008242-PA OS=Aedes aegypti GN=AAEL008242 PE=4 SV=1
CCG014767.1	3140	62	583	0.84844699	6.490660942	2.935468967	Up	0	0	Tetracycline resistance OS=Culex quinquefasciatus GN=CpipJ_CPIJ003321 PE=4 SV=1
CCG016129.1	2189	153	583	3.00336656	9.310495824	1.632277434	Up	0	0	AAEL006795-PA OS=Aedes aegypti GN=CYP9J15 PE=3 SV=2
CCG020347.1	1539	160	583	4.46728885	13.242804	1.567737182	Up	4.24E-14	1.44E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009391 PE=4 SV=1
CCG024265.1	732	213	583	12.5034883	27.84245268	1.154955657	Up	3.57E-13	9.59E-13	AAEL008801-PA OS=Aedes aegypti GN=AAEL008801 PE=4 SV=1
CCG010246.1	1158	9	584	0.33396167	17.63008086	5.722212761	Up	1.09E-06	2.01E-06	AAEL000828-PA OS=Aedes aegypti GN=AAEL000828 PE=3 SV=1
CCG008559.1	490	0	585	0.001	41.73590186	15.34900133	Up	4.44E-16	1.63E-15	Predicted protein OS=Nematostella vectensis GN=v1g229551 PE=3 SV=1
CCG021583.1	1989	38	585	0.82094013	10.28184611	3.646678505	Up	4.26E-14	1.45E-13	Actin OS=Culex quinquefasciatus GN=CpipJ_CPIJ003226 PE=3 SV=1
CCG021053.1	1632	89	585	2.34332499	12.53099995	2.418872587	Up	0	0	AAEL005019-PA OS=Aedes aegypti GN=AAEL005019 PE=4 SV=1
CCG010831.1	1872	137	585	3.14468677	10.92446149	1.796573935	Up	2.90E-13	8.03E-13	AAEL001282-PA OS=Aedes aegypti GN=AAEL001282 PE=4 SV=1
CCG023535.1	765	39	586	2.19061392	26.77849698	3.611667839	Up	7.33E-14	2.38E-13	AAEL000141-PA OS=Aedes aegypti GN=AAEL000141 PE=4 SV=1
CCG028121.1	2115	42	586	0.85329969	9.685839333	3.504752635	Up	7.66E-14	2.47E-13	AAEL000900-PA OS=Aedes aegypti GN=AAEL000900 PE=4 SV=1
CCG027441.1	1013	43	586	1.82398676	20.22265566	3.470805304	Up	8.46E-14	2.71E-13	AAEL000128-PA (Fragment) OS=Aedes aegypti GN=AAEL000128 PE=4 SV=1
CCG023065.1	1217	87	586	3.07178875	16.83282678	2.454126562	Up	0	0	AAEL004333-PA OS=Aedes aegypti GN=AAEL004333 PE=4 SV=1
CCG007941.1	1011	237	586	10.0730238	20.26266092	1.008326809	Up	2.32E-13	6.61E-13	AAEL009183-PA OS=Aedes aegypti GN=AAEL009183 PE=3 SV=1

CCG001094.1	631	2485	586	169.223123	32.46521424	-2.381960078	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007377 PE=4 SV=1
CCG027161.1	2013	37	587	0.78980635	10.19399328	3.690076531	Up	5.17E-14	1.74E-13	AAEL012059-PA OS=Aedes aegypti GN=AAEL012059 PE=4 SV=1
CCG005518.1	1523	982	587	27.7060272	13.47374161	-1.040049317	Down	4.13E-45	1.88E-44	AAEL000186-PA OS=Aedes aegypti GN=AAEL006186 PE=4 SV=1
CCG027169.1	1185	61	588	2.21194414	17.34638544	2.971248211	Up	1.75E-14	6.18E-14	AAEL001587-PA OS=Aedes aegypti GN=AAEL001587 PE=4 SV=1
CCG006308.1	1423	69	588	2.08356408	14.44516285	2.793461092	Up	0	0	CXAX prenyl protease z OS=Culex quinquefasciatus GN=CpipJ_CPIJ002690 PE=4 SV=1
CCG020129.1	1462	71	588	2.0867655	14.05982677	2.752238429	Up	0	0	AAEL012330-PA OS=Aedes aegypti GN=AAEL012330 PE=3 SV=1
CCG022951.1	642	90	588	6.02379457	32.01786097	2.410132452	Up	0	0	AAEL004945-PA (Fragment) OS=Aedes aegypti GN=AAEL004945 PE=4 SV=1
CCG021085.1	2013	1383	588	29.5216806	10.21135953	-1.531599892	Down	5.77E-114	3.12E-113	AAEL001645-PA OS=Aedes aegypti GN=AAEL001645 PE=4 SV=1
CCG014928.1	1299	99	589	3.27482966	15.85098154	2.275080408	Up	0	0	AAEL013546-PA OS=Aedes aegypti GN=AAEL013546 PE=3 SV=1
CCG014507.1	1146	141	589	5.28685216	17.96721206	1.764885675	Up	0	0	AAEL007219-PA OS=Aedes aegypti GN=AAEL007219 PE=4 SV=1
CCG022389.1	1657	1599	589	41.4656642	12.42632771	-1.738517196	Down	1.10E-157	6.33E-157	AAEL000505-PA OS=Aedes aegypti GN=AAEL000505 PE=4 SV=1
CCG004419.1	1046	2	590	0.0821601	19.71833967	7.906884348	Up	9.36E-05	0.00015327	Similar to Aedes aegypti 34 kDa salivary secreted protein 34k-2 OS=Aedes albopictus PE=2 SV=1
CCG006164.1	1695	18	590	0.45631577	12.16836773	4.736959347	Up	1.62E-09	3.32E-09	AAEL004698-PA OS=Aedes aegypti GN=AAEL004698 PE=4 SV=1
CCG015632.1	1414	35	590	1.06360729	14.58655113	3.777601331	Up	6.39E-14	2.11E-13	AAEL000874-PA OS=Aedes aegypti GN=AAEL008724 PE=4 SV=1
CCG002763.1	616	36	590	2.51121826	33.48276509	3.736959347	Up	6.13E-14	2.03E-13	-
CCG002254.1	1692	44	590	1.11741627	12.18994285	3.447452729	Up	3.91E-14	1.34E-13	AAEL000116-PA OS=Aedes aegypti GN=AAEL008116 PE=4 SV=1

CCG008533.1	1924	105	590	2.3450219	10.72005369	2.19263883	Up	0	0	AAEL010896-PA OS=Aedes aegypti GN=AAEL010896 PE=4 SV=1
CCG021008.1	1339	106	590	3.40163694	15.40357229	2.178963894	Up	0	0	AAEL003870-PA OS=Aedes aegypti GN=AAEL003870 PE=4 SV=1
CCG007641.1	13256	178	590	0.57699251	1.55592813	1.431150917	Up	1.15E-13	3.58E-13	AAEL001257-PA OS=Aedes aegypti GN=AAEL001257 PE=3 SV=1
CCG000326.1	576	4	591	0.29840093	35.86864856	6.909327524	Up	2.95E-05	5.04E-05	AAEL011582-PA OS=Aedes aegypti GN=AAEL011582 PE=4 SV=1
CCG017426.1	1260	6	591	0.20461778	16.39709649	6.324365023	Up	8.38E-06	1.48E-05	Putative salivary serpin OS=Aedes albopictus PE=2 SV=1
CCG003541.1	2043	128	591	2.69218112	10.11274673	1.909327524	Up	4.15E-14	1.42E-13	AAEL002080-PA OS=Aedes aegypti GN=AAEL002080 PE=4 SV=1
CCG008371.1	1227	129	591	4.51760046	16.83809419	1.898100269	Up	0	0	Stoucho protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009226 PE=4 SV=1
CCG016289.1	3157	153	591	2.08247368	6.544295715	1.651939681	Up	0	0	AAEL013725-PA OS=Aedes aegypti GN=AAEL013725 PE=4 SV=1
CCG027913.1	1110	220	591	8.51652398	18.61292034	1.12796781	Up	4.61E-13	1.20E-12	AAEL003409-PA OS=Aedes aegypti GN=AAEL003409 PE=3 SV=1
CCG024582.1	2631	1145	591	18.7002456	7.852657382	-1.251804359	Down	5.97E-70	2.96E-69	AGAP005258-PA OS=Anopheles gambiae GN=AGAP005258 PE=4 SV=4
CCG001403.1	1650	49	592	1.27607091	12.54260597	3.297056725	Up	0	0	AAEL012038-PA OS=Aedes aegypti GN=AAEL012038 PE=4 SV=1
CCG026257.1	2097	62	592	1.27044518	9.869003266	2.957570259	Up	0	0	AAEL001160-PA OS=Aedes aegypti GN=AAEL001160 PE=4 SV=1
CCG018478.1	3636	93	592	1.09906087	5.691776636	2.372607758	Up	4.04E-14	1.39E-13	AAEL007150-PA OS=Aedes aegypti GN=AAEL007150 PE=4 SV=1
CCG015229.1	1242	107	592	3.70190145	16.66288233	2.170299583	Up	0	0	AAEL004873-PA OS=Aedes aegypti GN=AAEL004873 PE=2 SV=1
CCG008217.1	831	151	592	7.80797825	24.90409127	1.67336183	Up	0	0	Proteasome subunit alpha type OS=Aedes aegypti GN=AAEL015007 PE=3 SV=1

CCG014320.1	1729	1916	592	47.6171264	11.96951987	-1.992115276	Down	4.02E-227	2.44E-226	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005148 PE=4 SV=1
CCG009500.1	1878	54	593	1.23555147	11.03847611	3.159313996	Up	0	0	AAEL012142-PA OS=Aedes aegypti GN=AAEL012142 PE=4 SV=1
CCG019022.1	2092	78	593	1.60212204	9.909301207	2.62879928	Up	0	0	AAEL013355-PA OS=Aedes aegypti GN=AAEL013355 PE=4 SV=1
CCG026212.1	3096	198	593	2.74806442	6.695819808	1.284844878	Up	0	0	AAEL007171-PA OS=Aedes aegypti GN=AAEL007171 PE=3 SV=1
CCG028116.1	2265	1428	593	27.0908525	9.152431844	-1.565578766	Down	1.95E-121	1.07E-120	Alpha-amylase 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017521 PE=3 SV=1
CCG007027.1	7467	176	594	1.01281282	2.780931619	1.457200706	Up	0	0	AAEL005044-PA OS=Aedes aegypti GN=AAEL005044 PE=4 SV=1
CCG010255.1	2112	190	594	3.86564847	9.832015342	1.346776716	Up	2.69E-13	7.52E-13	AAEL002910-PA OS=Aedes aegypti GN=AAEL002910 PE=4 SV=1
CCG011049.1	9529	1214	594	5.47436854	2.17916008	-1.328920382	Down	3.77E-81	1.92E-80	-
CCG016480.1	2298	36	595	0.67315511	9.051425013	3.749134061	Up	6.13E-14	2.03E-13	AAEL006655-PA OS=Aedes aegypti GN=AAEL006655 PE=4 SV=1
CCG000242.1	9048	152	595	0.72186115	2.298869881	1.671131549	Up	8.39E-14	2.69E-13	-
CCG001789.1	1472	163	595	4.75819751	14.13055345	1.570330908	Up	1.80E-13	5.25E-13	AAEL002772-PA OS=Aedes aegypti GN=AAEL002772 PE=4 SV=1
CCG026846.1	3115	206	595	2.84165821	6.677423653	1.232558535	Up	1.10E-13	3.44E-13	-
CCG011082.1	823	239	595	12.4784527	25.27360228	1.018192254	Up	6.72E-13	1.70E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017674 PE=4 SV=1
CCG027697.1	595	4	596	0.28887217	35.01703018	6.921481724	Up	2.95E-05	5.03E-05	histone H1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ012396 PE=3 SV=1
CCG010655.1	2069	108	596	2.24298276	10.07014643	2.166594222	Up	0	0	AAEL010613-PA OS=Aedes aegypti GN=AAEL010613 PE=4 SV=1
CCG002891.1	1635	118	596	3.10117962	12.74320058	2.038838675	Up	0	0	AGAP001363-PA OS=Anopheles gambiae GN=AgaP_AGAP001363 PE=4 SV=4

CCG008498.1	3709	128	596	1.48291346	5.61745294	1.921481724	Up	4.15E-14	1.42E-13	AAEL007256-PA OS=Aedes aegypti GN=AAEL007256 PE=4 SV=1
CCG014876.1	1326	128	596	4.14790802	15.71276995	1.921481724	Up	4.15E-14	1.42E-13	AAEL014308-PA OS=Aedes aegypti GN=AAEL014308 PE=4 SV=1
CCG002920.1	657	137	596	8.96020341	31.71253114	1.823449641	Up	2.90E-13	8.04E-13	AAEL001875-PA OS=Aedes aegypti GN=AAEL001875 PE=4 SV=2
CCG001313.1	2066	164	596	3.41095667	10.0847691	1.56392972	Up	1.14E-13	3.54E-13	AAEL000867-PA (Fragment) OS=Aedes aegypti GN=AAEL000867 PE=4 SV=1
CCG004397.1	682	208	596	13.105139	30.55004832	1.221042006	Up	3.93E-13	1.05E-12	AAEL001622-PA OS=Aedes aegypti GN=AAEL001622 PE=3 SV=1
CCG003223.1	1815	37	597	0.87596704	11.49867286	3.714446959	Up	5.17E-14	1.74E-13	AAEL008320-PA OS=Aedes aegypti GN=AAEL008320 PE=4 SV=1
CCG024461.1	2285	49	597	0.92145164	9.133519139	3.309190481	Up	0	0	AGAP003794-PA OS=Anopheles gambiae GN=AgaP_AGAP003794 PE=4 SV=5
CCG023521.1	1188	75	597	2.71273577	17.56741686	2.695081635	Up	0	0	tRNA-dihydrouridine synthase OS=Aedes aegypti GN=AAEL005898 PE=3 SV=1
CCG000927.1	684	111	598	6.97315869	30.56293788	2.131899012	Up	0	0	AAEL013309-PA OS=Aedes aegypti GN=AAEL013309 PE=4 SV=1
CCG013050.1	2373	130	598	2.3540099	8.809544673	1.903947065	Up	0	0	AAEL009427-PA OS=Aedes aegypti GN=AAEL009427 PE=4 SV=1
CCG004448.2	1764	145	598	3.5320927	11.8509351	1.746405788	Up	0	0	AAEL001426-PA OS=Aedes aegypti GN=AAEL001426 PE=4 SV=1
CCG013032.1	620	215	598	14.9007951	33.71782179	1.178122028	Up	0	0	AAEL000722-PA OS=Aedes aegypti GN=AAEL000722 PE=4 SV=1
CCG001579.1	5457	1670	598	13.1499829	3.830868519	-1.779317509	Down	5.10E-170	2.96E-169	Strinthe OS=Culex quinquefasciatus GN=Cuq1_Cuq1014866 PE=4 SV=1
CCG002942.1	2022	22	599	0.46752431	10.35608694	4.469293778	Up	8.38E-11	1.80E-10	AAEL007457-PA OS=Aedes aegypti GN=AAEL007457 PE=4 SV=1
CCG006675.1	1003	169	599	7.24016466	20.87737566	1.52784596	Up	0	0	AAEL007280-PA OS=Aedes aegypti GN=AAEL007280 PE=4 SV=1
CCG000392.1	3825	1701	599	19.1088937	5.47451184	-1.803442029	Down	2.10E-176	1.23E-175	AAEL010379-PA OS=Aedes aegypti GN=AAEL010379 PE=3 SV=1

CCG009860.1	1473	1778	599	51.8670659	14.21589123	-1.867314212	Down	2.50E-193	1.48E-192	AAEL012404-PA OS=Aedes aegypti GN=AAEL012464 PE=3 SV=1
CCG002643.1	915	2149	599	100.920175	22.88525441	-2.140724371	Down	1.51E-279	9.42E-279	AAEL010163-PA OS=Aedes aegypti GN=AAEL010163 PE=4 SV=1
CCG003865.1	1071	2435	599	97.6949615	19.551828	-2.32098066	Down	0	0	AAEL002457-PA OS=Aedes aegypti GN=AAEL002457 PE=4 SV=1
CCG013787.1	1549	74	600	2.05278267	13.54097228	2.721678529	Up	0	0	AAEL005325-PA OS=Aedes aegypti GN=AAEL005325 PE=4 SV=1
CCG022629.1	2927	75	600	1.10103522	7.16602872	2.702313204	Up	0	0	AAEL009308-PA OS=Aedes aegypti GN=AAEL009308 PE=4 SV=1
CCG004160.1	1821	107	600	2.52485536	11.51837785	2.189664908	Up	0	0	AAEL004343-PA OS=Aedes aegypti GN=AAEL004343 PE=3 SV=1
CCG024270.1	3834	175	600	1.9613207	5.470778838	1.479920782	Up	3.55E-13	9.59E-13	AAEL008035-PA OS=Aedes aegypti GN=AAEL008035 PE=4 SV=1
CCG009926.1	1023	9	601	0.37803286	20.53756045	5.763609383	Up	1.09E-06	2.01E-06	AAEL010910-PA OS=Aedes aegypti GN=AAEL010910 PE=4 SV=1
CCG026426.1	1854	145	601	3.36063189	11.33221378	1.753625295	Up	0	0	AAEL001151-PA OS=Aedes aegypti GN=AAEL001151 PE=3 SV=1
CCG003830.1	6285	1513	601	10.3441859	3.342867835	-1.629661888	Down	3.20E-136	1.79E-135	AAEL009888-PA OS=Aedes aegypti GN=AAEL009888 PE=4 SV=1
CCG002231.1	1227	2933	601	102.714125	17.12300272	-2.584626971	Down	0	0	AAEL013431-PA (Fragment) OS=Aedes aegypti GN=AAEL013431 PE=4 SV=1
CCG006577.1	1890	51	602	1.15950078	11.13485853	3.263507539	Up	0	0	AAEL006381-PA OS=Aedes aegypti GN=AAEL006381 PE=4 SV=1
CCG022814.1	1722	66	602	1.64692363	12.22118619	2.891538761	Up	0	0	AAEL004352-PA OS=Aedes aegypti GN=AAEL004352 PE=4 SV=1
CCG024056.1	2226	70	602	1.35124952	9.454125165	2.806649864	Up	0	0	AAEL013014-PA OS=Aedes aegypti GN=AAEL013014 PE=4 SV=1
CCG009177.1	2379	72	602	1.30047116	8.846104505	2.766007879	Up	0	0	Receptor protein tyrosine kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ000058 PE=3 SV=1
CCG017968.1	918	157	602	7.34885439	22.92470873	1.641312132	Up	0	0	AAEL008373-PA OS=Aedes aegypti GN=AAEL008373 PE=4 SV=1

CCG017266.1	1242	206	602	7.12702523	16.94434993	1.249432353	Up	1.10E-13	3.43E-13	AAEL006868-PA OS=Aedes aegypti GN=AAEL006868 PE=3 SV=1
CCG000205.1	4136	226535	602	2353.51761	5.088221136	-8.85344156	Down	0	0	ADP,ATP carrier protein 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005941 PE=3 SV=1
CCG004332.1	684	86	603	5.40262745	30.81848084	2.512062641	Up	0	0	Single-stranded DNA-binding protein OS=Aedes aegypti GN=AAEL000147 PE=3 SV=1
CCG026604.1	2835	183	603	2.77370774	7.43556998	1.422627557	Up	3.05E-13	8.42E-13	AAEL003865-PA OS=Aedes aegypti GN=AAEL003865 PE=4 SV=1
CCG021820.1	1407	198	603	6.04691361	14.98211862	1.308970776	Up	0	0	AAEL010563-PA OS=Aedes aegypti GN=AAEL010563 PE=3 SV=1
CCG027447.1	2477	228	603	3.95522789	8.510230478	1.105437382	Up	1.05E-13	3.28E-13	AAEL013769-PA OS=Aedes aegypti GN=AAEL013769 PE=4 SV=1
CCG027462.2	1722	11	604	0.27448727	12.26178814	5.481286325	Up	2.67E-07	5.03E-07	AAEL011836-PA OS=Aedes aegypti GN=AAEL011836 PE=4 SV=1
CCG021644.2	1548	161	604	4.46907447	13.64005114	1.609801065	Up	0	0	AAEL000403-PA OS=Aedes aegypti GN=AAEL000403 PE=4 SV=1
CCG010168.1	6087	983	604	6.93925565	3.468835086	-1.000329663	Down	1.93E-42	8.70E-42	AAEL005871-PA OS=Aedes aegypti GN=AAEL005871 PE=4 SV=1
CCG017201.1	2307	1366	604	25.4428511	9.152492055	-1.475023825	Down	1.60E-106	8.55E-106	Phosphoserine aminotransferase OS=Aedes aegypti PE=2 SV=1
CCG017023.1	891	61	605	2.94181124	23.73710151	3.012367198	Up	1.75E-14	6.17E-14	AAEL014296-PA OS=Aedes aegypti GN=AAEL014296 PE=4 SV=1
CCG020064.1	2064	83	605	1.7279496	10.24697551	2.568065105	Up	1.32E-13	4.02E-13	AAEL014145-PA OS=Aedes aegypti GN=AAEL014145 PE=4 SV=1
CCG002801.1	1302	99	605	3.26728397	16.24405334	2.313747916	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011327 PE=4 SV=1
CCG023328.1	1005	102	605	4.36110739	21.04453477	2.270679194	Up	0	0	AAEL005203-PA OS=Aedes aegypti GN=AAEL005203 PE=4 SV=1
CCG027760.1	1053	68	606	2.77487365	20.11843848	2.858024346	Up	0	0	KRR1 small subunit processome component OS=Aedes aegypti GN=AAEL000708 PE=3 SV=1

CCG022229.1	540	98	606	7.7982111	39.23095504	2.330777343	Up	0	0	AGAP010252-PA OS=Anopheles gambiae GN=AGAP010252 PE=3 SV=3
CCG007221.2	2013	111	606	2.36941905	10.52395217	2.151071321	Up	0	0	AAEL005074-PA OS=Aedes aegypti GN=AAEL005074 PE=4 SV=1
CCG019571.1	2965	153	606	2.21732526	7.144929418	1.688099345	Up	0	0	AAEL001756-PA OS=Aedes aegypti GN=AAEL001756 PE=3 SV=1
CCG012442.1	912	188	606	8.85779617	23.22885496	1.390898336	Up	1.80E-13	5.24E-13	AAEL011988-PA OS=Aedes aegypti GN=AAEL011988 PE=4 SV=1
CCG014493.1	1803	233	606	5.55293853	11.74970367	1.081301043	Up	0	0	AAEL006343-PA OS=Aedes aegypti GN=AAEL006343 PE=4 SV=1
CCG017615.1	1733	30	607	0.74384999	12.24447432	4.040975314	Up	3.71E-13	9.93E-13	AGAP013226-PA OS=Anopheles gambiae GN=AgaP_AGAP013226 PE=4 SV=1
CCG023635.1	1065	67	607	2.7032603	19.92457653	2.88177672	Up	0	0	AAEL015082-PA (Fragment) OS=Aedes aegypti GN=AAEL015082 PE=4 SV=1
CCG001035.1	690	151	607	9.40352163	30.75315073	1.709461171	Up	0	0	-
CCG010591.1	1643	1320	607	34.5222457	12.91520024	-1.418456304	Down	3.53E-97	1.86E-96	AAEL010151-PA OS=Aedes aegypti GN=CYP6N16 PE=3 SV=2
CCG013097.1	1029	25	608	1.04396829	20.65561932	4.306384527	Up	9.32E-12	2.08E-11	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1
CCG017890.1	2160	45	608	0.8952028	9.840107536	3.458387621	Up	0	0	AAEL013593-PA OS=Aedes aegypti GN=AAEL013593 PE=4 SV=1
CCG008817.1	1734	102	608	2.52763145	12.2575734	2.277815375	Up	0	0	AAEL007244-PA OS=Aedes aegypti GN=AAEL007244 PE=4 SV=1
CCG009595.1	900	215	609	10.2649922	23.65510062	1.204418772	Up	0	0	AAEL010592-PA OS=Aedes aegypti GN=AAEL010592 PE=4 SV=1
CCG000118.1	1047	1318	609	54.0917958	20.33389738	-1.411523033	Down	2.52E-96	1.32E-95	AAEL006526-PA OS=Aedes aegypti GN=AAEL006526 PE=4 SV=1
CCG003021.2	1536	77	610	2.15408175	13.88316981	2.688192096	Up	0	0	AAEL010805-PA OS=Aedes aegypti GN=AAEL010805 PE=4 SV=1
CCG020316.1	1563	183	611	5.0310054	13.66571152	1.441641935	Up	3.05E-13	8.42E-13	AAEL004757-PA OS=Aedes aegypti GN=AAEL004757 PE=4 SV=1

CCG020992.2	1940	218	611	4.82855781	11.01005521	1.189157449	Up	0	0	AAEL001893-PA OS=Aedes aegypti GN=AAEL001893 PE=4 SV=1
CCG028549.1	2356	240	611	4.37722254	9.066004715	1.050451178	Up	0	0	AAEL009174-PA OS=Aedes aegypti GN=AAEL009174 PE=2 SV=1
CCG026999.1	630	241	611	16.4376286	33.90397954	1.044452437	Up	2.37E-13	6.72E-13	Ubiquitin carboxyl-terminal hydrolase (Fragment) OS=Anopheles gambiae GN=AGAP009348 PE=3 SV=4
CCG013805.1	1137	1182	611	44.6703837	18.78584618	-1.249672547	Down	5.81E-72	2.89E-71	AAEL012311-PA OS=Aedes aegypti GN=AAEL012311 PE=3 SV=1
CCG007431.1	2460	1286	611	22.4630401	8.682726467	-1.371333154	Down	4.88E-90	2.54E-89	AAEL004090-PA (Fragment) OS=Aedes aegypti GN=AAEL004090 PE=4 SV=1
CCG027427.1	1665	159	612	4.1034161	12.84952876	1.646818091	Up	5.53E-14	1.85E-13	AAEL011371-PA OS=Aedes aegypti GN=AAEL011371 PE=4 SV=1
CCG003976.1	1869	187	612	4.29927254	11.44701198	1.412806587	Up	3.31E-13	9.04E-13	AAEL009736-PA OS=Aedes aegypti GN=AAEL009736 PE=4 SV=1
CCG027301.2	1194	20	613	0.71976105	17.94759101	4.640128373	Up	3.69E-10	7.70E-10	AAEL004128-PA OS=Aedes aegypti GN=AAEL004128 PE=4 SV=1
CCG002816.1	1525	51	613	1.43702063	14.05208109	3.289631126	Up	0	0	AAEL011361-PA OS=Aedes aegypti GN=AAEL011361 PE=4 SV=1
CCG014300.1	1236	71	613	2.46832618	17.33772141	2.812309348	Up	0	0	AAEL003572-PA OS=Aedes aegypti GN=AAEL003572 PE=4 SV=1
CCG005066.1	2219	73	613	1.41360551	9.657243651	2.772231909	Up	0	0	AAEL002864-PA OS=Aedes aegypti GN=AAEL002864 PE=4 SV=1
CCG004176.1	1868	176	613	4.04854031	11.47185421	1.502624849	Up	0	0	AAEL007797-PA OS=Aedes aegypti GN=AAEL007797 PE=4 SV=1
CCG020178.1	1362	2574	613	81.2071196	15.73379123	-2.367739871	Down	0	0	AAEL014137-PA OS=Aedes aegypti GN=CLIPB25 PE=3 SV=1
CCG010120.1	885	0	614	0.001	24.25353891	14.56590765	Up	4.44E-16	1.64E-15	AAEL014276-PA OS=Aedes aegypti GN=AAEL014276 PE=4 SV=1
CCG005939.1	701	2	614	0.12259553	30.6196604	7.964408049	Up	9.36E-05	0.00015403	-
CCG017158.1	705	78	614	4.7540983	30.44593183	2.67900583	Up	0	0	AAEL005028-PA OS=Aedes aegypti GN=AAEL005028 PE=4 SV=1

CCG014675.1	1887	95	614	2.16328818	11.37487119	2.394552441	Up	0	0	AAEL010795-PA OS=Aedes aegypti GN=AAEL010795 PE=4 SV=1
CCG010360.1	1802	1066	614	25.419388	11.91142172	-1.093583674	Down	7.33E-53	3.46E-52	AAEL001548-PA OS=Aedes aegypti GN=AAEL001548 PE=4 SV=1
CCG020389.1	4542	1532	614	14.4935344	4.725755601	-1.616792533	Down	2.41E-136	1.35E-135	AAEL007316-PA OS=Aedes aegypti GN=AAEL007316 PE=4 SV=1
CCG023245.1	795	7352	614	397.375458	26.99922256	-3.879513002	Down	0	0	NADH dehydrogenase iron-sulfur protein 3, mitochondrial OS=Culex quinquefasciatus GN=CpipJ_CPIJ013503 PE=3 SV=1
CCG022858.1	1820	68	615	1.60546261	11.81282429	2.879292963	Up	0	0	Predicted acetyltransferase OS=Aedes aegypti PE=2 SV=1
CCG006588.2	1314	176	615	5.75545913	16.36175054	1.507324185	Up	0	0	AAEL008932-PA OS=Aedes aegypti GN=AAEL008932 PE=3 SV=1
CCG022511.1	5013	205	615	1.75719042	4.288717378	1.287275705	Up	1.83E-13	5.31E-13	Canalicular multispecific organic anion transporter 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ016918 PE=3 SV=1
CCG001424.1	763	32	616	1.80213828	28.22319593	3.969099745	Up	1.77E-13	5.16E-13	AAEL009555-PA OS=Aedes aegypti GN=AAEL009555 PE=4 SV=1
CCG004751.1	6676	120	616	0.7723739	3.225628893	2.062209149	Up	0	0	AAEL012178-PA OS=Aedes aegypti GN=AAEL012178 PE=4 SV=1
CCG007242.1	1375	203	616	6.34389536	15.66130799	1.303763827	Up	9.28E-14	2.93E-13	AAEL013643-PA OS=Aedes aegypti GN=AAEL013643 PE=4 SV=1
CCG013204.5	2007	237	616	5.07415401	10.72959566	1.080356496	Up	2.32E-13	6.62E-13	AAEL007865-PA OS=Aedes aegypti GN=AAEL007865 PE=4 SV=1
CCG002091.1	822	1796	616	93.8852109	26.19744342	-1.84147189	Down	1.41E-191	8.38E-191	AAEL011315-PA OS=Aedes aegypti GN=AAEL011315 PE=4 SV=1
CCG001439.5	2036	43	617	0.90751404	10.59393751	3.545175128	Up	8.46E-14	2.70E-13	AAEL003261-PA OS=Aedes aegypti GN=AAEL003261 PE=4 SV=1
CCG001062.1	723	138	617	8.20169208	29.83299691	1.862915426	Up	1.49E-13	4.45E-13	AAEL010753-PA OS=Aedes aegypti GN=AAEL010753 PE=4 SV=1
CCG009830.1	1428	159	617	4.78444524	15.10452155	1.658556928	Up	5.53E-14	1.85E-13	AAEL007967-PA OS=Aedes aegypti GN=AAEL007967 PE=4 SV=1

CCG000052.1	1569	183	617	5.01176637	13.74713625	1.455740045	Up	3.05E-13	8.41E-13	AAEL014628-PA OS=Aedes aegypti GN=AAEL014628 PE=4 SV=1
CCG004008.1	1191	239	617	8.62280988	18.11020719	1.070573075	Up	6.72E-13	1.70E-12	AAEL008416-PA OS=Aedes aegypti GN=AAEL008416 PE=4 SV=1
CCG028029.1	1055	1712	617	69.7290859	20.44479315	-1.770027103	Down	5.59E-173	3.25E-172	AAEL000431-PA OS=Aedes aegypti GN=AAEL000431 PE=4 SV=1
CCG006203.1	992	11834	617	512.604677	21.74320239	-4.559210297	Down	0	0	AAEL006070-PA OS=Aedes aegypti GN=AAEL006070 PE=3 SV=1
CCG014788.1	477	26	618	2.34216583	45.29185544	4.273336514	Up	4.60E-12	1.05E-11	AAEL013145-PA OS=Aedes aegypti GN=AAEL013145 PE=4 SV=1
CCG010131.1	621	58	618	4.01327634	34.78939621	3.115795237	Up	0	0	AAEL003944-PA OS=Aedes aegypti GN=AAEL003944 PE=4 SV=1
CCG011134.1	1338	146	618	4.68877523	16.14664802	1.783951673	Up	9.33E-15	3.34E-14	SV1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017451 PE=4 SV=1
CCG007830.1	747	167	618	9.60635299	28.92130528	1.590071939	Up	1.62E-13	4.77E-13	AAEL013467-PA OS=Aedes aegypti GN=AAEL013467 PE=4 SV=1
CCG015665.1	3261	24	619	0.3162446	6.635747722	4.391146302	Up	1.93E-11	4.23E-11	Predicted protein (Fragment) OS=Nematostella vectensis GN=v1g103876 PE=4 SV=1
CCG025259.1	1258	31	619	1.05887263	17.20125065	4.021912493	Up	2.40E-13	6.76E-13	AAEL007461-PA OS=Aedes aegypti GN=AAEL007461 PE=4 SV=1
CCG010600.1	1566	89	619	2.44208581	13.81811834	2.500375372	Up	0	0	AAEL001827-PA OS=Aedes aegypti GN=AAEL001827 PE=4 SV=1
CCG011843.1	2136	183	619	3.6813958	10.13069912	1.460408965	Up	3.05E-13	8.42E-13	AAEL012938-PA OS=Aedes aegypti GN=AAEL012938 PE=4 SV=1
CCG027887.1	1840	1497	619	34.9596156	11.76042028	-1.571749703	Down	6.12E-128	3.39E-127	AAEL013434-PA OS=Aedes aegypti GN=AAEL013434 PE=4 SV=1
CCG012828.1	1629	40	620	1.05511933	13.30517594	3.656509514	Up	6.86E-14	2.24E-13	Ubiquitin carboxyl-terminal hydrolase (Fragment) OS=Aedes aegypti GN=AAEL003517 PE=3 SV=1
CCG022593.1	1521	50	620	1.4125488	14.24992216	3.334581419	Up	0	0	AAEL002784-PA OS=Aedes aegypti GN=AAEL002784 PE=4 SV=1
CCG024952.1	3570	1014	620	12.204849	6.071185322	-1.007404328	Down	3.21E-44	1.46E-43	AAEL003026-PA OS=Aedes aegypti GN=AAEL003026 PE=4 SV=1

CCG015336.1	815	1149	620	60.5794173	26.5940265	-1.187725474	Down	1.05E-64	5.14E-64	Putative uncharacterized protein OS=Aedes aegypti PE=2 SV=1
CCG001267.1	1683	24	622	0.61275914	12.91981471	4.398121473	Up	1.93E-11	4.23E-11	AAEL009218-PA OS=Aedes aegypti GN=AAEL009218 PE=3 SV=1
CCG020950.1	571	30	622	2.25760427	38.08064475	4.076193378	Up	3.71E-13	9.94E-13	-
CCG017451.1	1668	112	622	2.88525796	13.03600009	2.175729052	Up	0	0	AAEL013204-PA OS=Aedes aegypti GN=AAEL013204 PE=4 SV=1
CCG011269.1	882	1893	622	92.2241583	24.65311582	-1.903374722	Down	4.02E-211	2.41E-210	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC013386 PE=4 SV=1
CCG016245.1	3037	3151	622	44.5826914	7.159712925	-2.638510065	Down	0	0	Transporter OS=Aedes aegypti GN=AAEL010681 PE=3 SV=1
CCG002715.1	738	128	623	7.4527453	29.51084882	1.985401557	Up	4.15E-14	1.42E-13	AAEL009060-PA OS=Aedes aegypti GN=AAEL009060 PE=4 SV=1
CCG022255.1	1380	131	623	4.07901104	15.78188872	1.951978555	Up	1.26E-13	3.87E-13	Glycylpeptide N- tetradecanoyltransferase OS=Aedes aegypti GN=AAEL005092 PE=3 SV=1
CCG016630.1	1207	154	623	5.48246821	18.04391585	1.718615016	Up	1.82E-13	5.29E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_05216 PE=4 SV=1
CCG022308.1	1141	157	623	5.91257523	19.08764805	1.690780808	Up	0	0	AAEL008615-PA OS=Aedes aegypti GN=AAEL008615 PE=4 SV=1
CCG023823.1	1533	172	623	4.82113135	14.20678828	1.559136802	Up	1.35E-13	4.09E-13	AAEL015082-PA (Fragment) OS=Aedes aegypti GN=AAEL015082 PE=4 SV=1
CCG003249.1	1194	208	623	7.48551491	18.24037389	1.284961839	Up	3.93E-13	1.05E-12	AAEL006435-PA OS=Aedes aegypti GN=AAEL006435 PE=3 SV=1
CCG009921.1	959	94	624	4.21184051	22.74657425	2.433126571	Up	0	0	-
CCG018898.1	2067	132	624	2.74407594	10.55344204	1.943321303	Up	1.46E-13	4.36E-13	AAEL004791-PA OS=Aedes aegypti GN=AAEL004791 PE=4 SV=1
CCG018668.1	3942	136	624	1.48246674	5.533730265	1.900252581	Up	1.70E-13	4.98E-13	AAEL003495-PA (Fragment) OS=Aedes aegypti GN=AAEL003495 PE=4 SV=1

CCG024421.3	1860	148	624	3.41909716	11.72793801	1.778262057	Up	0	0	AAEL003546-PA OS=Aedes aegypti GN=AAEL003546 PE=4 SV=1
CCG013718.1	2979	155	624	2.23575323	7.322579626	1.711591017	Up	3.42E-13	9.27E-13	AAEL005025-PA OS=Aedes aegypti GN=AAEL005025 PE=4 SV=1
CCG016326.1	2067	2329	624	48.4163096	10.55344204	-2.197779502	Down	0	0	SLC12-like K,Cl cotransporter OS=Aedes aegypti GN=KCC1A PE=2 SV=1
CCG025779.1	1128	50	625	1.90468682	19.36961257	3.346169394	Up	0	0	Nucleoprotein N OS=Tibrogargan virus GN=N PE=4 SV=1
CCG021997.1	1779	114	625	2.75354117	12.28157559	2.157135569	Up	8.48E-14	2.70E-13	AAEL005677-PA OS=Aedes aegypti GN=AAEL005677 PE=4 SV=1
CCG025940.1	2031	137	625	2.89850007	10.75771688	1.8919935	Up	2.90E-13	8.04E-13	AAEL011309-PA OS=Aedes aegypti GN=AAEL011309 PE=3 SV=1
CCG008709.1	2364	1613	625	29.3190279	9.242353208	-1.66550514	Down	1.10E-149	6.27E-149	Mis transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ017440 PE=4 SV=1
CCG024662.1	813	17	626	0.89850614	26.91744312	4.904869209	Up	3.39E-09	6.88E-09	AGAP013324-PA OS=Anopheles gambiae GN=AgaP_AGAP013324 PE=4 SV=1
CCG003761.1	694	85	626	5.26286375	31.53297011	2.582941115	Up	0	0	AAEL003435-PA OS=Aedes aegypti GN=AAEL003435 PE=4 SV=1
CCG023052.1	761	126	626	7.11456841	28.75674279	2.015052127	Up	1.34E-13	4.06E-13	AAEL000163-PA OS=Aedes aegypti GN=AAEL000163 PE=4 SV=1
CCG012807.1	2588	136	626	2.25806952	8.45590466	1.904869209	Up	1.70E-13	4.98E-13	AG2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ015325 PE=3 SV=1
CCG003529.1	1425	151	626	4.55328416	15.35710966	1.753927311	Up	0	0	AAEL009331-PA OS=Aedes aegypti GN=AAEL009331 PE=4 SV=1
CCG025636.2	1602	158	626	4.23796384	13.66035035	1.688551303	Up	0	0	AAEL008468-PA OS=Aedes aegypti GN=AAEL008468 PE=4 SV=1
CCG011149.1	675	221	626	14.0686094	32.42056483	1.204429491	Up	0	0	AAEL007349-PA OS=Aedes aegypti GN=AAEL007349 PE=4 SV=1
CCG020584.1	2208	1063	626	20.6869692	9.911178107	-1.061593831	Down	2.68E-50	1.25E-49	AAEL002412-PA OS=Aedes aegypti GN=AAEL002412 PE=4 SV=1
CCG002983.1	1193	53	627	1.90896558	18.37287472	3.266714382	Up	0	0	AAEL007708-PA OS=Aedes aegypti GN=AAEL007708 PE=4 SV=1

CCG007406.1	1911	60	628	1.34912825	11.48812026	3.090043357	Up	0	0	AAEL004049-PA OS=Aedes aegypti GN=AAEL004049 PE=4 SV=1
CCG011912.1	1341	65	628	2.08279847	16.37121388	2.97456614	Up	0	0	DNA replication licensing factor Mcm6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009047 PE=3 SV=1
CCG019425.1	1128	77	628	2.9332177	19.46258671	2.730147412	Up	0	0	AAEL003385-PA OS=Aedes aegypti GN=AAEL003385 PE=4 SV=1
CCG009386.1	1282	86	628	2.8825251	17.12464728	2.570669198	Up	0	0	AAEL003456-PA OS=Aedes aegypti GN=AAEL003456 PE=4 SV=1
CCG028043.1	2799	112	628	1.71940346	7.843443306	2.189579031	Up	0	0	AAEL000924-PA OS=Aedes aegypti GN=AAEL000924 PE=4 SV=1
CCG008685.1	465	209	628	19.3132786	47.21246842	1.289574821	Up	0	0	Beta-arrestin 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017114 PE=4 SV=1
CCG013832.1	1125	175	629	6.68418094	19.54556097	1.548018299	Up	3.55E-13	9.57E-13	Proteasome subunit beta type OS=Aedes aegypti GN=AAEL002401 PE=3 SV=1
CCG020450.1	1368	247	629	7.7584243	16.07365211	1.050862179	Up	0	0	AAEL004186-PA (Fragment) OS=Aedes aegypti GN=AAEL004186 PE=4 SV=1
CCG023854.1	1407	48	630	1.46591845	15.65295975	3.416558721	Up	5.35E-14	1.79E-13	AAEL008550-PA OS=Aedes aegypti GN=AAEL008550 PE=4 SV=1
CCG012732.1	1599	70	630	1.88110158	13.77342987	2.872238205	Up	0	0	AAEL008867-PA OS=Aedes aegypti GN=AAEL008867 PE=4 SV=1
CCG002660.1	2583	79	630	1.31421178	8.526408969	2.697740474	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002374 PE=4 SV=1
CCG010813.1	2528	150	630	2.54962824	8.711912329	1.772702532	Up	0	0	AAEL009242-PA OS=Aedes aegypti GN=AAEL009242 PE=4 SV=1
CCG025220.1	2805	223	630	3.4161322	7.851591575	1.200621322	Up	0	0	AAEL005811-PA OS=Aedes aegypti GN=AAEL005811 PE=4 SV=1
CCG000848.1	546	30	631	2.36097443	40.40049935	4.096918803	Up	3.71E-13	9.93E-13	AAEL005263-PA OS=Aedes aegypti GN=AAEL005263 PE=4 SV=1
CCG018883.1	1356	45	631	1.42598677	16.2674577	3.511956302	Up	0	0	AAEL011423-PA OS=Aedes aegypti GN=AAEL011423 PE=4 SV=1

CCG011261.1	1668	61	631	1.57143514	13.22462389	3.073072061	Up	1.75E-14	6.17E-14	AAEL013204-PA OS=Aedes aegypti GN=AAEL013204 PE=4 SV=1
CCG027242.1	859	71	631	3.55163115	25.67947921	2.854062279	Up	0	0	AAEL006838-PA OS=Aedes aegypti GN=AAEL006838 PE=4 SV=1
CCG018763.1	1452	144	631	4.26146128	15.19192331	1.833884397	Up	7.53E-14	2.43E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004597 PE=4 SV=1
CCG013928.1	732	221	631	12.9731029	30.13479869	1.215906839	Up	0	0	AAEL000488-PA OS=Aedes aegypti GN=AAEL000488 PE=4 SV=1
CCG013160.1	1440	34	632	1.01456318	15.34279925	3.918631111	Up	8.82E-14	2.80E-13	AAEL004055-PA OS=Aedes aegypti GN=AAEL004055 PE=4 SV=1
CCG026841.1	1105	39	632	1.51657887	19.99423613	3.720691733	Up	7.33E-14	2.38E-13	AAEL003648-PA OS=Aedes aegypti GN=AAEL003648 PE=4 SV=1
CCG027165.1	1813	41	632	0.97173697	12.18622776	3.648541947	Up	1.02E-13	3.19E-13	AAEL009187-PA OS=Aedes aegypti GN=AAEL009187 PE=4 SV=1
CCG025096.1	4041	47	632	0.49977172	5.467367216	3.4515051	Up	0	0	AAEL014493-PA OS=Aedes aegypti GN=AAEL014493 PE=4 SV=1
CCG005024.1	996	52	632	2.2433998	22.18236036	3.305654234	Up	0	0	AAEL013114-PA OS=Aedes aegypti GN=AAEL013114 PE=4 SV=1
CCG023235.1	897	218	632	10.4430347	24.63058074	1.237909627	Up	0	0	AAEL007593-PA OS=Aedes aegypti GN=CLIPC2 PE=3 SV=1
CCG019184.1	856	131	633	6.57597574	25.8511556	1.974951892	Up	1.26E-13	3.87E-13	Condensin complex subunit 2 OS=Aedes aegypti GN=AAEL005962 PE=3 SV=1
CCG003150.1	1191	134	633	4.83454613	18.5798398	1.942285703	Up	0	0	AAEL007279-PA OS=Aedes aegypti GN=AAEL007279 PE=3 SV=1
CCG025035.1	1375	155	633	4.84386099	16.09351942	1.732250488	Up	3.42E-13	9.26E-13	AAEL011526-PA OS=Aedes aegypti GN=GPRMTH5 PE=4 SV=2
CCG020483.1	305	157	633	22.118847	72.55275147	1.713754144	Up	0	0	-
CCG001273.1	3242	235	633	3.11470933	6.825598148	1.131857947	Up	4.34E-13	1.14E-12	AAEL008115-PA OS=Aedes aegypti GN=AAEL008115 PE=4 SV=1
CCG017552.1	4965	1591	633	13.769355	4.456916253	-1.627343227	Down	7.52E-143	4.24E-142	-
CCG027939.1	2134	1874	633	37.7344342	10.36953571	-1.863530344	Down	4.28E-203	2.56E-202	AAEL012291-PA OS=Aedes aegypti GN=AAEL012291 PE=4 SV=1

CCG003055.1	1566	23	634	0.63110083	14.15296774	4.487090278	Up	4.01E-11	8.66E-11	AAEL000422-PA OS=Aedes aegypti GN=AAEL000422 PE=4 SV=1
CCG023750.1	1914	53	634	1.18986203	11.57970087	3.282731779	Up	0	0	AAEL009254-PA OS=Aedes aegypti GN=AAEL009254 PE=3 SV=1
CCG011608.1	2055	97	634	2.02825511	10.78518125	2.410739392	Up	3.62E-14	1.25E-13	AAEL013546-PA OS=Aedes aegypti GN=AAEL013548 PE=4 SV=1
CCG026570.1	1466	106	634	3.10695216	15.11838163	2.282731779	Up	0	0	AAEL013521-PA OS=Aedes aegypti GN=AAEL013521 PE=3 SV=1
CCG022913.1	1551	127	634	3.51847601	14.28984363	2.021967547	Up	1.57E-13	4.67E-13	Spire OS=Culex quinquefasciatus GN=CpipJ_CPIJ011319 PE=4 SV=1
CCG010318.1	2041	146	634	3.07377817	10.85916094	1.820827675	Up	9.33E-15	3.34E-14	AGAP002258-PB OS=Anopheles gambiae GN=AgaP_AGAP002258 PE=4 SV=1
CCG020490.1	828	47	635	2.43910329	26.80978955	3.458337134	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009765 PE=4 SV=1
CCG027947.2	1362	81	635	2.5554688	16.29846237	2.673075983	Up	4.46E-14	1.51E-13	RNA methyltransferase OS=Culex quinquefasciatus GN=CpipJ_CPIJ009249 PE=4 SV=1
CCG019001.1	816	150	635	7.89884827	27.20405116	1.784107295	Up	0	0	AAEL014671-PA OS=Aedes aegypti GN=AAEL014671 PE=4 SV=1
CCG016968.1	798	153	635	8.23855814	27.81767638	1.755538143	Up	0	0	AAEL000789-PA OS=Aedes aegypti GN=AAEL000789 PE=4 SV=1
CCG024760.1	1734	50	636	1.23903502	12.82206691	3.371339969	Up	0	0	AAEL000445-PA OS=Aedes aegypti GN=AAEL006445 PE=4 SV=1
CCG002998.1	1311	67	636	2.19601237	16.95916402	2.949106969	Up	0	0	AAEL007316-PA OS=Aedes aegypti GN=AAEL007318 PE=4 SV=1
CCG025097.1	3286	111	636	1.45150351	6.766118085	2.220780293	Up	0	0	AAEL014497-PA OS=Aedes aegypti GN=AAEL014497 PE=4 SV=1
CCG026038.1	4266	116	636	1.16842223	5.211782472	2.157215164	Up	1.16E-13	3.58E-13	CK3F complex subunit 3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000292 PE=4 SV=1
CCG004688.1	1518	176	636	4.98199822	14.64655074	1.55576454	Up	0	0	AAEL004892-PA OS=Aedes aegypti GN=AAEL004892 PE=4 SV=1

CCG025016.1	429	187	636	18.7303971	51.82625647	1.468301699	Up	3.31E-13	9.04E-13	AAEL002261-PA OS=Aedes aegypti GN=AAEL002261 PE=3 SV=1
CCG000394.1	1182	203	636	7.37974292	18.81003725	1.349860242	Up	9.28E-14	2.93E-13	AAEL010388-PA OS=Aedes aegypti GN=AAEL010388 PE=4 SV=1
CCG017285.1	2754	216	636	3.37017526	8.073153242	1.260308657	Up	1.38E-13	4.17E-13	AAEL003227-PA OS=Aedes aegypti GN=AAEL003227 PE=4 SV=1
CCG003789.1	1464	13	637	0.38156185	15.21067097	5.317023048	Up	6.34E-08	1.23E-07	AAEL009404-PA OS=Aedes aegypti GN=AAEL009404 PE=4 SV=1
CCG013984.1	3146	146	637	1.99414534	7.078328768	1.827638207	Up	9.33E-15	3.33E-14	AAEL004237-PA OS=Aedes aegypti GN=AAEL004237 PE=4 SV=1
CCG023264.1	1787	191	637	4.59273604	12.46134432	1.440033938	Up	6.68E-14	2.19E-13	Transmembrane protein 15 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002894 PE=4 SV=1
CCG013066.1	1541	88	638	2.45382002	14.47331641	2.560294199	Up	2.93E-14	1.02E-13	AAEL001580-PA OS=Aedes aegypti GN=AAEL001580 PE=4 SV=1
CCG009749.1	939	161	638	7.36754768	23.75226899	1.688808939	Up	0	0	AAEL006727-PA OS=Aedes aegypti GN=AAEL006727 PE=4 SV=1
CCG007246.2	726	2612	638	154.596345	30.72090989	-2.331213364	Down	0	0	AAEL006711-PA OS=Aedes aegypti GN=AAEL006711 PE=4 SV=1
CCG025085.1	939	20	639	0.91522331	23.78949825	4.70005723	Up	3.69E-10	7.72E-10	Sub1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009759 PE=4 SV=1
CCG021426.1	1572	27	639	0.73802979	14.21013922	4.267097823	Up	2.28E-12	5.41E-12	AAEL013154-PA OS=Aedes aegypti GN=AAEL013154 PE=4 SV=1
CCG021732.1	1369	114	639	3.57819558	16.31726724	2.189095311	Up	8.48E-14	2.70E-13	AAEL000465-PA OS=Aedes aegypti GN=AAEL000465 PE=4 SV=1
CCG004503.1	652	140	639	9.22663013	34.26125592	1.892702308	Up	0	0	AAEL007790-PA OS=Aedes aegypti GN=AAEL007790 PE=4 SV=1
CCG024277.1	2617	165	639	2.70921139	8.535857416	1.655663111	Up	1.31E-14	4.67E-14	AAEL014407-PA OS=Aedes aegypti GN=AAEL014407 PE=4 SV=1
CCG027864.2	1657	192	639	4.97899158	13.48119424	1.437022824	Up	0	0	AAEL012390-PA OS=Aedes aegypti GN=AAEL012390 PE=4 SV=1
CCG017486.1	954	9	640	0.40537485	23.45209343	5.854316297	Up	1.09E-06	2.01E-06	Similar to Aedes aegypti 34 kDa salivary secreted protein 34k-2 OS=Aedes albopictus PE=2 SV=1

CCG021431.1	2894	74	640	1.09874235	7.730925064	2.814787933	Up	0	0	AAEL000707-PA OS=Aedes aegypti GN=AAEL000707 PE=4 SV=1
CCG011681.1	1140	157	640	5.9177617	19.62569924	1.72962055	Up	0	0	AAEL000013-PA OS=Aedes aegypti GN=AAEL000013 PE=4 SV=1
CCG002019.1	1320	12	641	0.39063395	16.97595107	5.44153125	Up	1.30E-07	2.49E-07	AAEL010808-PA OS=Aedes aegypti GN=AAEL010808 PE=4 SV=1
CCG005547.1	1509	27	641	0.76884217	14.84973851	4.271606248	Up	2.28E-12	5.40E-12	AAEL000260-PA OS=Aedes aegypti GN=AAEL000260 PE=4 SV=1
CCG012150.1	1899	33	641	0.74670945	11.80002918	3.982099631	Up	1.31E-13	4.00E-13	AAEL012142-PA OS=Aedes aegypti GN=AAEL012142 PE=4 SV=1
CCG007518.1	732	41	641	2.40677475	30.61237078	3.668941746	Up	1.02E-13	3.20E-13	Eukaryotic translation initiation factor 6 OS=Aedes aegypti GN=EIF6 PE=3 SV=1
CCG014864.1	1605	66	641	1.76697974	13.96152985	2.982099631	Up	0	0	AAEL008536-PA OS=Aedes aegypti GN=AAEL008536 PE=4 SV=1
CCG024498.1	1942	99	641	2.19052715	11.5387515	2.39713713	Up	0	0	AAEL004892-PA OS=Aedes aegypti GN=AAEL004892 PE=4 SV=1
CCG000443.1	1329	106	641	3.42723241	16.86098978	2.298573296	Up	0	0	AAEL000846-PA OS=Aedes aegypti GN=AAEL000846 PE=4 SV=1
CCG015967.1	2694	126	641	2.00972033	8.31783794	2.049213827	Up	1.34E-13	4.06E-13	AAEL007618-PA OS=Aedes aegypti GN=AAEL007618 PE=4 SV=1
CCG011074.1	2376	182	641	3.29145274	9.431083927	1.51869911	Up	0	0	AAEL013900-PA (Fragment) OS=Aedes aegypti GN=AAEL013900 PE=4 SV=1
CCG000751.1	1728	237	641	5.89341846	12.9677404	1.137750502	Up	2.32E-13	6.61E-13	AAEL007399-PA OS=Aedes aegypti GN=AAEL007399 PE=4 SV=1
CCG012516.1	1272	1089	641	36.7877681	17.616553	-1.062294488	Down	1.54E-51	7.22E-51	AAEL008793-PA OS=Aedes aegypti GN=AAEL008793 PE=4 SV=1
CCG007604.1	5748	2148	641	16.0575835	3.898443878	-2.042284528	Down	7.25E-263	4.49E-262	AAEL002171-PA (Fragment) OS=Aedes aegypti GN=AAEL002171 PE=4 SV=1
CCG003955.1	741	41	642	2.37754267	30.28773777	3.671190686	Up	1.02E-13	3.19E-13	AAEL009212-PA OS=Aedes aegypti GN=AAEL009212 PE=4 SV=1

CCG011531.1	1708	125	642	3.14474053	13.14005485	2.062958406	Up	2.35E-14	8.23E-14	Phosphatidylinositol N-acetylglucosaminyl transferase subunit Q OS=Culex quinquefasciatus GN=CpipJ_CPIJ005945 PE=4 SV=1
CCG002813.1	2193	19	643	0.3722868	10.24996442	4.783060618	Up	7.73E-10	1.60E-09	Zinc finger protein 558 OS=Culex quinquefasciatus GN=CpipJ_CPIJ018144 PE=4 SV=1
CCG002447.1	852	77	643	3.88341498	26.38283094	2.76420159	Up	0	0	AAEL011057-PA OS=Aedes aegypti GN=AAEL011057 PE=4 SV=1
CCG005029.1	1614	243	643	6.46942101	13.92699626	1.106175628	Up	1.53E-14	5.41E-14	Peptidoglycan recognition protein LE OS=Armigeres subalbus GN=PGRP-LE PE=2 SV=1
CCG000016.1	3513	1228	643	15.0204481	6.398568735	-1.231106714	Down	5.53E-73	2.76E-72	AAEL008859-PA OS=Aedes aegypti GN=AAEL008859 PE=3 SV=1
CCG005629.1	1398	99	644	3.04292112	16.10381276	2.403873462	Up	0	0	AGAP001363-PA OS=Anopheles gambiae GN=AgaP_AGAP001363 PE=4 SV=4
CCG005803.1	1287	143	644	4.77441496	17.49271969	1.873358745	Up	0	0	AAEL002282-PA OS=Aedes aegypti GN=AAEL002282 PE=4 SV=1
CCG003153.1	1108	170	644	6.59282932	20.3187096	1.623839146	Up	3.53E-13	9.53E-13	AAEL004376-PA OS=Aedes aegypti GN=AAEL004376 PE=4 SV=1
CCG027955.1	1494	187	644	5.37840721	15.06902961	1.486335622	Up	3.31E-13	9.04E-13	AAEL002819-PA OS=Aedes aegypti GN=AAEL002819 PE=4 SV=1
CCG009081.1	759	30	645	1.69840848	29.70762651	4.128577959	Up	3.71E-13	9.93E-13	AAEL006683-PA OS=Aedes aegypti GN=AAEL006683 PE=4 SV=1
CCG012282.1	591	55	645	3.99887547	38.15243404	3.254108841	Up	0	0	AAEL011205-PA OS=Aedes aegypti GN=AAEL011205 PE=4 SV=1
CCG021312.1	1662	2	646	0.05170847	13.58787412	8.037703559	Up	9.36E-05	0.00015326	AAEL000889-PA (Fragment) OS=Aedes aegypti GN=AAEL000889 PE=4 SV=2
CCG026688.1	1821	135	646	3.18556517	12.40145348	1.960887961	Up	1.90E-13	5.48E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006732 PE=4 SV=1

CCG014868.1	2787	232	646	3.57695674	8.102994903	1.179722563	Up	2.91E-13	8.05E-13	AAEL014406-PA OS=Aedes aegypti GN=AAEL014406 PE=4 SV=1
CCG027569.1	897	31	647	1.4850187	25.2151673	4.085738795	Up	2.40E-13	6.78E-13	AAEL002496-PA OS=Aedes aegypti GN=AAEL002496 PE=4 SV=1
CCG013233.1	630	72	647	4.91082681	35.90159535	2.870010104	Up	0	0	AAEL010760-PA OS=Aedes aegypti GN=AAEL010760 PE=4 SV=1
CCG013461.1	1707	118	647	2.97037416	13.25014943	2.157292056	Up	0	0	AAEL006289-PA (Fragment) OS=Aedes aegypti GN=AAEL006289 PE=4 SV=1
CCG015333.1	1368	116	648	3.64363247	16.55918373	2.184182212	Up	1.16E-13	3.58E-13	AAEL004131-PA OS=Aedes aegypti GN=AAEL004131 PE=4 SV=1
CCG017859.1	1416	232	649	7.040239	16.02254352	1.186406877	Up	2.91E-13	8.06E-13	AAEL009958-PA OS=Aedes aegypti GN=AAEL009958 PE=4 SV=1
CCG022697.1	13984	1477	649	4.53849385	1.622420025	-1.484066239	Down	4.56E-116	2.48E-115	AAEL006136-PA OS=Aedes aegypti GN=AAEL006136 PE=3 SV=1
CCG013994.1	1002	4	650	0.17153587	22.67752485	7.046609112	Up	2.95E-05	5.03E-05	Uncharacterized protein OS=Tetrapisispora phaffii (strain ATCC 24235 / CBS 4417 / NBRC 1672 / NRRL Y-8282 / UCD 70-5) GN=TPHA0B00750 PE=4 SV=1
CCG014461.1	1062	136	650	5.50271554	21.39630876	1.95914627	Up	1.70E-13	4.99E-13	AAEL013291-PA OS=Aedes aegypti GN=AAEL013291 PE=4 SV=1
CCG026443.1	1113	1193	650	46.0583049	20.41588491	-1.173769216	Down	6.93E-66	3.39E-65	AAEL005238-PA OS=Aedes aegypti GN=AAEL005238 PE=4 SV=1
CCG011199.1	2715	1407	650	22.2682934	8.369384863	-1.411797502	Down	9.88E-103	5.25E-102	Uncharacterized protein OS=Anopheles darlingi GN=AND_15753 PE=4 SV=1
CCG017376.1	1275	140	651	4.71824537	17.84928485	1.91954392	Up	0	0	AAEL010427-PA OS=Aedes aegypti GN=AAEL010427 PE=4 SV=1
CCG026947.1	1566	193	651	5.29575912	14.53246372	1.4563699	Up	6.08E-14	2.02E-13	AAEL001157-PA OS=Aedes aegypti GN=AAEL001157 PE=4 SV=1
CCG011110.1	3861	3456	651	38.4624198	5.894285983	-2.706060565	Down	0	0	AAEL004824-PA (Fragment) OS=Aedes aegypti GN=AAEL004824 PE=4 SV=1

CCG018467.1	984	23	652	1.00437388	23.16341103	4.527479402	Up	4.01E-11	8.67E-11	AAEL012018-PA OS=Aedes aegypti GN=AAEL012018 PE=4 SV=1
CCG016823.1	2214	50	652	0.97040954	10.29484935	3.407185168	Up	0	0	AAEL001598-PA OS=Aedes aegypti GN=AAEL001598 PE=4 SV=1
CCG022434.1	2445	139	652	2.44286017	9.322207139	1.932100285	Up	1.46E-13	4.37E-13	AAEL010729-PA (Fragment) OS=Aedes aegypti GN=AAEL010729 PE=4 SV=1
CCG003482.1	1563	147	652	4.04129942	14.58272326	1.851369013	Up	7.02E-14	2.29E-13	AAEL004057-PA OS=Aedes aegypti GN=AAEL004057 PE=4 SV=1
CCG013698.1	2718	1540	652	24.3463544	8.385870661	-1.537673278	Down	2.03E-127	1.12E-126	AAEL005464-PA OS=Aedes aegypti GN=AAEL005464 PE=4 SV=1
CCG007283.1	6267	2049	652	14.0489846	3.636954916	-1.949662911	Down	7.20E-236	4.39E-235	AAEL000239-PA OS=Aedes aegypti GN=AAEL000239 PE=4 SV=1
CCG008179.1	980	51	653	2.23618007	23.29362728	3.380827043	Up	0	0	AAEL006294-PA OS=Aedes aegypti GN=AAEL006294 PE=4 SV=1
CCG017059.1	1506	59	653	1.68340926	15.15787167	3.170609336	Up	0	0	AAEL008788-PA OS=Aedes aegypti GN=AAEL008788 PE=4 SV=1
CCG007338.1	1362	193	653	6.08895652	16.76046603	1.460795348	Up	6.08E-14	2.02E-13	AAEL007460-PA OS=Aedes aegypti GN=AAEL007460 PE=4 SV=1
CCG015056.1	693	215	653	13.3311586	32.94048302	1.305059536	Up	0	0	AAEL013774-PA OS=Aedes aegypti GN=AAEL013774 PE=4 SV=1
CCG008177.1	1852	38	654	0.88166842	12.34487743	3.807532516	Up	4.26E-14	1.45E-13	AAEL005783-PA OS=Aedes aegypti GN=AAEL005783 PE=4 SV=1
CCG009097.1	1620	73	654	1.93629051	14.11278581	2.86563547	Up	0	0	AAEL014483-PA OS=Aedes aegypti GN=AAEL014483 PE=4 SV=1
CCG011692.1	1693	78	654	1.97970425	13.50426049	2.77005781	Up	0	0	AGAP008634-PA (Fragment) OS=Anopheles gambiae GN=AGAP008634 PE=4 SV=4
CCG010046.1	714	147	654	8.84671007	32.02060646	1.855787684	Up	7.02E-14	2.29E-13	AAEL012195-PA OS=Aedes aegypti GN=AAEL012195 PE=4 SV=1
CCG026752.1	1128	148	654	5.63787298	20.2683626	1.846006664	Up	0	0	AAEL013389-PA OS=Aedes aegypti GN=AAEL013389 PE=4 SV=1
CCG017449.1	1177	247	654	9.01743793	19.424565	1.107092798	Up	0	0	AAEL013207-PA OS=Aedes aegypti GN=AAEL013207 PE=4 SV=1

CCG007340.1	20582	1078	654	2.25057691	1.110811049	-1.018681433	Down	9.65E-48	4.46E-47	AAEL009055-PA OS=Aedes aegypti GN=AAEL009055 PE=4 SV=1
CCG024833.1	1734	44	655	1.09035082	13.20511608	3.598232682	Up	3.91E-14	1.34E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000812 PE=4 SV=1
CCG021817.1	1140	52	655	1.96002298	20.08567657	3.357224582	Up	0	0	Cell growth-regulating nuclear protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013225 PE=4 SV=1
CCG008420.1	1551	95	655	2.63193088	14.76316653	2.487808692	Up	0	0	AAEL012734-PA OS=Aedes aegypti GN=AAEL012734 PE=4 SV=1
CCG014222.1	2374	113	655	2.04531593	9.645185883	2.237485338	Up	1.42E-13	4.26E-13	AAEL012593-PA (Fragment) OS=Aedes aegypti GN=AAEL012593 PE=4 SV=1
CCG017383.1	1297	123	655	4.07500182	17.65433407	2.115149795	Up	0	0	AAEL002801-PA OS=Aedes aegypti GN=AAEL002801 PE=4 SV=1
CCG011656.2	3932	106	656	1.15839061	5.832306603	2.331944754	Up	0	0	AAEL000396-PA OS=Aedes aegypti GN=AAEL000396 PE=4 SV=1
CCG024937.1	2190	191	656	3.74758873	10.47152035	1.48243638	Up	6.68E-14	2.19E-13	AAEL000965-PA OS=Aedes aegypti GN=AAEL000965 PE=4 SV=1
CCG025738.1	2730	200	656	3.14796591	8.400230609	1.416009019	Up	0	0	AAEL000962-PA OS=Aedes aegypti GN=AAEL000962 PE=4 SV=1
CCG007698.1	1144	225	656	8.45121529	20.04600486	1.246084017	Up	0	0	AAEL011189-PA OS=Aedes aegypti GN=AAEL011189 PE=4 SV=1
CCG027695.1	506	2	657	0.16984085	45.3904898	8.062062764	Up	9.36E-05	0.00015342	Predicted protein OS=Nematostella vectensis GN=v1g229551 PE=3 SV=1
CCG021056.1	1085	69	657	2.7326375	21.16828372	2.953538307	Up	0	0	AAEL006637-PA OS=Aedes aegypti GN=AAEL006637 PE=4 SV=1
CCG015000.1	1935	131	657	2.90906214	11.86955444	2.028639763	Up	1.26E-13	3.87E-13	AAEL002006-PA OS=Aedes aegypti GN=AAEL002006 PE=4 SV=1
CCG021019.1	568	193	657	14.6006317	40.43589408	1.469605727	Up	6.08E-14	2.02E-13	26S proteasome regulatory complex subunit RPN5/PSMD12 OS=Aedes aegypti GN=AAEL014325 PE=2 SV=1

CCG007395.1	718	1176	657	70.3793982	31.9882839	-1.137609581	Down	7.27E-62	3.53E-61	AAEL004071-PA OS=Aedes aegypti GN=AAEL004071 PE=4 SV=1
CCG006258.1	792	0	658	0.001	29.04361883	14.8259336	Up	4.44E-16	1.63E-15	AAEL008580-PA OS=Aedes aegypti GN=AAEL008580 PE=4 SV=1
CCG027921.1	1641	169	658	4.42528041	14.01739556	1.663377541	Up	0	0	Leucokinin receptor protein OS=Aedes aegypti PE=2 SV=1
CCG027239.1	915	173	658	8.12433234	25.13939466	1.62962875	Up	3.60E-13	9.69E-13	AAEL007688-PA OS=Aedes aegypti GN=AAEL007688 PE=4 SV=1
CCG020856.1	1849	176	658	4.0901424	12.44053332	1.604825359	Up	0	0	ShfC domain-containing histone demethylation protein 3C OS=Culex quinquefasciatus GN=CpipJ_CPIJ011492 PE=4 SV=1
CCG001234.1	816	208	658	10.9530696	28.18939475	1.363817259	Up	3.93E-13	1.05E-12	AAEL009686-PA (Fragment) OS=Aedes aegypti GN=AAEL009686 PE=4 SV=1
CCG017745.1	1531	217	658	6.0904196	15.02452392	1.302705745	Up	1.47E-13	4.37E-13	AAEL010842-PA OS=Aedes aegypti GN=AAEL010842 PE=4 SV=1
CCG028580.1	2130	219	658	4.41801497	10.79931743	1.289469918	Up	4.32E-13	1.14E-12	AAEL011389-PA OS=Aedes aegypti GN=AAEL011389 PE=4 SV=1
CCG027182.1	1080	136	659	5.41100362	21.33102259	1.978985018	Up	1.70E-13	4.98E-13	AAEL008932-PA OS=Aedes aegypti GN=AAEL008932 PE=3 SV=1
CCG004862.2	4211	143	659	1.45919545	5.470791829	1.906576522	Up	0	0	AAEL001410-PA (Fragment) OS=Aedes aegypti GN=AAEL001410 PE=4 SV=1
CCG011338.1	2559	165	659	2.77061595	9.002541771	1.700125645	Up	1.31E-14	4.67E-14	AAEL010280-PA OS=Aedes aegypti GN=AAEL010280 PE=4 SV=1
CCG013650.1	981	213	659	9.32982005	23.48369459	1.331738239	Up	3.57E-13	9.60E-13	AAEL003961-PA OS=Aedes aegypti GN=AAEL003961 PE=3 SV=1
CCG010792.1	909	259	659	12.2433017	25.34378921	1.049639571	Up	0	0	AAEL006735-PA OS=Aedes aegypti GN=AAEL006735 PE=2 SV=1
CCG013100.1	1739	45	660	1.11192528	13.26766111	3.576782322	Up	0	0	AAEL005286-PA OS=Aedes aegypti GN=AAEL005286 PE=4 SV=1
CCG012639.3	1624	234	660	6.19145191	14.20718145	1.198270698	Up	4.60E-13	1.20E-12	AAEL004971-PA OS=Aedes aegypti GN=AAEL004971 PE=4 SV=1

CCG010695.1	1821	20	661	0.47193558	12.68941293	4.74889157	Up	3.69E-10	7.71E-10	AAEL007176-PA OS=Aedes aegypti GN=AAEL007176 PE=3 SV=1
CCG013550.1	1592	29	661	0.78274014	14.51471165	4.21283867	Up	6.66E-13	1.69E-12	AAEL000261-PA OS=Aedes aegypti GN=AAEL000261 PE=4 SV=1
CCG026930.1	1893	61	661	1.384656	12.20677282	3.140082328	Up	1.75E-14	6.18E-14	AAEL010730-PA OS=Aedes aegypti GN=AAEL010730 PE=4 SV=1
CCG018019.1	1580	95	661	2.58362328	14.62494997	2.500964057	Up	0	0	tRNA pseudouridine synthase OS=Aedes aegypti GN=AAEL007354 PE=3 SV=1
CCG022838.1	1401	69	662	2.11628243	16.51847197	2.964476154	Up	0	0	AAEL012147-PA OS=Aedes aegypti GN=AAEL012147 PE=4 SV=1
CCG010373.1	1004	9935	662	425.203499	23.05017851	-4.205303642	Down	0	0	AAEL006070-PA OS=Aedes aegypti GN=AAEL006070 PE=3 SV=1
CCG007095.1	264	0	663	0.001	87.79294508	16.42181739	Up	4.44E-16	1.65E-15	-
CCG008932.1	1083	34	663	1.34900367	21.40105032	3.987715423	Up	8.82E-14	2.80E-13	AAEL011245-PA OS=Aedes aegypti GN=AAEL011245 PE=4 SV=1
CCG001460.1	807	35	663	1.86361922	28.72036865	3.945895247	Up	6.39E-14	2.11E-13	AAEL009995-PA OS=Aedes aegypti GN=AAEL009995 PE=4 SV=1
CCG015546.1	1508	49	663	1.39623143	15.3695872	3.46046842	Up	0	0	AAEL009023-PA OS=Aedes aegypti GN=AAEL009023 PE=4 SV=1
CCG014584.1	636	63	663	4.25643598	36.44235456	3.09789834	Up	0	0	AAEL005256-PA OS=Aedes aegypti GN=AAEL005256 PE=4 SV=1
CCG021068.1	987	196	663	8.53299694	23.48261145	1.46046842	Up	7.93E-14	2.55E-13	AGAP002890-PA OS=Anopheles gambiae GN=AgaP_AGAP002890 PE=4 SV=5
CCG006014.1	984	225	663	9.82539663	23.55420478	1.261397073	Up	0	0	Short-chain dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ009166 PE=3 SV=1
CCG016080.1	1875	42	664	0.96252206	12.37989108	3.685035212	Up	7.66E-14	2.48E-13	AAEL011220-PA OS=Aedes aegypti GN=AAEL011220 PE=4 SV=1
CCG002496.1	1286	46	664	1.53702006	18.04999672	3.553790679	Up	3.40E-14	1.18E-13	AAEL001600-PA OS=Aedes aegypti GN=AAEL001600 PE=4 SV=1
CCG005365.1	546	99	664	7.79121562	42.51336223	2.447996015	Up	0	0	AAEL014307-PA OS=Aedes aegypti GN=AAEL014307 PE=4 SV=1

CCG023213.1	2073	187	664	3.87618928	11.19744128	1.530458175	Up	3.31E-13	9.03E-13	AAEL000434-PA OS=Aedes aegypti GN=AAEL000434 PE=3 SV=1
CCG018688.1	3108	201	664	2.77893071	7.468563635	1.426300944	Up	0	0	AAEL014331-PA OS=Aedes aegypti GN=AAEL014331 PE=3 SV=1
CCG009461.1	1052	0	665	0.001	22.09815024	14.43163799	Up	4.44E-16	1.64E-15	AAEL015464-PA OS=Aedes aegypti GN=AAEL015464 PE=3 SV=1
CCG025700.1	2243	97	665	1.85825424	10.36435758	2.479610892	Up	3.62E-14	1.25E-13	AAEL013316-PA (Fragment) OS=Aedes aegypti GN=AAEL013316 PE=4 SV=1
CCG005587.1	507	185	665	15.6792917	45.8525721	1.548142274	Up	0	0	AAEL009995-PA OS=Aedes aegypti GN=AAEL009995 PE=4 SV=1
CCG001998.1	408	5	666	0.52658988	57.06424591	6.759763476	Up	1.59E-05	2.76E-05	Short form D7 salivary protein SD7-1 OS=Aedes albopictus PE=2 SV=1
CCG013549.1	1616	41	666	1.09019747	14.40730961	3.724139566	Up	1.02E-13	3.20E-13	AAEL000292-PA OS=Aedes aegypti GN=AAEL000292 PE=4 SV=1
CCG026888.1	1836	62	666	1.45104768	12.68094354	3.127495261	Up	0	0	AAEL000128-PA (Fragment) OS=Aedes aegypti GN=AAEL000128 PE=4 SV=1
CCG010533.1	750	0	667	0.001	31.08956081	14.92414262	Up	4.44E-16	1.65E-15	AAEL000499-PA OS=Aedes aegypti GN=SPZ1A PE=4 SV=1
CCG000408.1	670	4	667	0.25653573	34.80174717	7.083856155	Up	2.95E-05	5.03E-05	AAEL004562-PA OS=Aedes aegypti GN=AAEL004562 PE=4 SV=1
CCG013684.1	693	31	667	1.92216706	33.64671083	4.129659845	Up	2.40E-13	6.77E-13	AAEL013774-PA OS=Aedes aegypti GN=AAEL013774 PE=4 SV=1
CCG013194.1	1505	151	667	4.31124912	15.49313662	1.845451416	Up	0	0	AAEL012260-PA OS=Aedes aegypti GN=AAEL012260 PE=4 SV=1
CCG007774.1	1224	156	667	5.4765348	19.04997599	1.798453936	Up	1.42E-13	4.27E-13	AAEL009495-PA (Fragment) OS=Aedes aegypti GN=AAEL009495 PE=4 SV=1
CCG021156.1	2560	216	667	3.62557136	9.108269768	1.328968653	Up	1.38E-13	4.17E-13	AAEL007974-PA OS=Aedes aegypti GN=AAEL007974 PE=4 SV=1
CCG008789.1	3030	46	668	0.65234581	7.706973229	3.56245554	Up	3.40E-14	1.18E-13	AAEL008509-PA OS=Aedes aegypti GN=AAEL008509 PE=3 SV=1
CCG019789.1	2067	167	668	3.47167183	11.29759501	1.702313204	Up	1.62E-13	4.77E-13	AAEL000434-PA OS=Aedes aegypti GN=AAEL000434 PE=3 SV=1

CCG021242.1	1230	265	668	9.25770705	18.98547064	1.036168947	Up	4.94E-13	1.28E-12	AAEL001874-PA OS=Aedes aegypti GN=AAEL001874 PE=4 SV=1
CCG006000.1	641	0	669	0.001	36.48531538	15.1550283	Up	4.44E-16	1.65E-15	-
CCG017582.1	1158	65	669	2.41194538	20.1961029	3.065807791	Up	0	0	AAEL002106-PA OS=Aedes aegypti GN=AAEL002106 PE=4 SV=1
CCG017309.1	2283	199	669	3.7455003	10.2440154	1.451550984	Up	0	0	AAEL002345-PA OS=Aedes aegypti GN=AAEL002345 PE=4 SV=1
CCG003192.1	540	211	669	16.7900259	43.30942067	1.367076416	Up	1.30E-13	3.98E-13	AAEL000947-PA OS=Aedes aegypti GN=AAEL000947 PE=2 SV=1
CCG001827.1	1373	234	670	7.32331966	17.059028	1.21996577	Up	4.60E-13	1.20E-12	AAEL001945-PA OS=Aedes aegypti GN=AAEL001945 PE=4 SV=1
CCG000979.1	1071	24	671	0.96290722	21.90196425	4.507519659	Up	1.93E-11	4.23E-11	AAEL012418-PA OS=Aedes aegypti GN=AAEL012418 PE=4 SV=1
CCG010177.2	681	63	671	3.97517369	34.44493937	3.115202237	Up	0	0	AAEL006473-PA OS=Aedes aegypti GN=AAEL006473 PE=4 SV=1
CCG015444.1	582	106	671	7.82610287	40.30413009	2.364561705	Up	0	0	AAEL011841-PA OS=Aedes aegypti GN=AAEL011841 PE=3 SV=1
CCG011501.1	3310	231	671	2.99879417	7.086708071	1.240733119	Up	1.29E-13	3.97E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001501 PE=4 SV=1
CCG028033.1	678	1582	671	100.262714	34.59735061	-1.535051724	Down	1.63E-130	9.06E-130	AAEL005593-PA OS=Aedes aegypti GN=AAEL005593 PE=4 SV=1
CCG016495.1	597	4152	671	298.844787	39.29146351	-2.927108568	Down	0	0	-
CCG026259.1	783	49	672	2.68903831	30.00250574	3.479920782	Up	0	0	Proliferating cell nuclear antigen OS=Aedes aegypti GN=PCNA PE=2 SV=1
CCG012890.1	2695	119	672	1.8973649	8.716869013	2.199812863	Up	0	0	AAEL009500-PA OS=Aedes aegypti GN=AAEL009500 PE=4 SV=1
CCG028556.1	2796	125	672	1.92103606	8.401989267	2.128846342	Up	2.35E-14	8.22E-14	AAEL010497-PA OS=Aedes aegypti GN=AAEL010497 PE=4 SV=1
CCG011512.1	1468	160	672	4.68334982	16.0026989	1.772702532	Up	4.24E-14	1.45E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014088 PE=4 SV=1
CCG019649.1	849	158	673	7.99672328	27.71133129	1.79299515	Up	0	0	AAEL005401-PA OS=Aedes aegypti GN=AAEL005401 PE=4 SV=1

CCG023211.1	453	209	673	19.8248886	51.93580633	1.389416766	Up	0	0	Putative salivary secreted peptide OS=Aedes albopictus PE=2 SV=1
CCG002520.1	1269	110	674	3.72472089	18.56728018	2.317558271	Up	0	0	AAEL001492-PA OS=Aedes aegypti GN=AAEL001492 PE=4 SV=1
CCG014826.1	1092	215	674	8.46015837	21.57681185	1.350725135	Up	0	0	AAEL010247-PA OS=Aedes aegypti GN=AAEL010247 PE=4 SV=1
CCG000447.1	831	1111	674	57.4481049	28.35364446	-1.018725116	Down	3.75E-49	1.74E-48	AAEL000545-PA OS=Aedes aegypti GN=CTLMA11 PE=4 SV=1
CCG012596.1	651	43	675	2.83824668	36.24706117	3.674792141	Up	8.46E-14	2.71E-13	AAEL013066-PA (Fragment) OS=Aedes aegypti GN=AAEL013066 PE=4 SV=1
CCG023127.1	2612	1333	675	21.9290414	9.034011034	-1.279404169	Down	6.65E-84	3.40E-83	AAEL011863-PA (Fragment) OS=Aedes aegypti GN=AAEL011863 PE=4 SV=1
CCG024745.1	1030	15	676	0.62577283	22.94349039	5.196302044	Up	1.48E-08	2.92E-08	AAEL009880-PA OS=Aedes aegypti GN=AAEL009880 PE=4 SV=1
CCG019674.1	4149	156	676	1.61563717	5.695780935	1.817790421	Up	1.42E-13	4.27E-13	AAEL008592-PA OS=Aedes aegypti GN=AAEL008592 PE=3 SV=1
CCG018806.1	1556	166	676	4.58417477	15.18752898	1.728153209	Up	1.83E-13	5.32E-13	AAEL012390-PA OS=Aedes aegypti GN=AAEL012390 PE=4 SV=1
CCG018121.1	918	177	676	8.28501419	25.74269619	1.63558709	Up	0	0	AAEL001451-PA OS=Aedes aegypti GN=AAEL001451 PE=4 SV=1
CCG007137.1	1142	193	676	7.2619604	20.69334072	1.510735603	Up	6.08E-14	2.02E-13	-
CCG002047.2	1515	1592	676	45.1536749	15.59854462	-1.53343198	Down	4.00E-131	2.23E-130	AAEL009099-PA OS=Aedes aegypti GN=AAEL009099 PE=4 SV=1
CCG017694.1	2043	2837	677	59.6696706	11.58431394	-2.364825208	Down	0	0	AAEL007103-PA OS=Aedes aegypti GN=AAEL007103 PE=4 SV=1
CCG008317.1	1746	108	678	2.65792173	13.57486349	2.352567165	Up	0	0	AAEL013211-PA OS=Aedes aegypti GN=AAEL013211 PE=4 SV=1
CCG010619.1	2172	124	678	2.45315244	10.91239026	2.153258357	Up	1.61E-13	4.77E-13	AAEL009924-PA OS=Aedes aegypti GN=AAEL009924 PE=4 SV=1
CCG018768.2	2128	166	678	3.35196238	11.13802239	1.732415236	Up	1.83E-13	5.32E-13	AAEL011701-PA OS=Aedes aegypti GN=AAEL011701 PE=4 SV=1
CCG005464.1	930	230	678	10.6269236	25.48571145	1.261964616	Up	3.85E-13	1.03E-12	AAEL009110-PA OS=Aedes aegypti GN=AAEL009110 PE=4 SV=1

CCG009107.1	669	252	678	16.185909	35.42856749	1.130174743	Up	5.42E-14	1.81E-13	AAEL014029-PA OS=Aedes aegypti GN=AAEL014029 PE=3 SV=1
CCG003979.1	927	252	678	11.6810929	25.56818948	1.130174743	Up	5.42E-14	1.81E-13	AAEL009735-PA OS=Aedes aegypti GN=AAEL009735 PE=4 SV=1
CCG016063.1	1268	51	679	1.72827797	18.71977124	3.437155626	Up	0	0	AAEL004943-PA OS=Aedes aegypti GN=AAEL004943 PE=4 SV=1
CCG000193.1	1563	142	679	3.90384025	15.18660904	1.959833849	Up	2.62E-13	7.35E-13	AAEL008613-PA OS=Aedes aegypti GN=AAEL008613 PE=4 SV=1
CCG017030.1	603	183	679	13.0405662	39.36429507	1.59388113	Up	3.05E-13	8.41E-13	AAEL003774-PA OS=Aedes aegypti GN=AAEL003774 PE=4 SV=1
CCG019074.1	1176	0	680	0.001	20.21396956	14.30306504	Up	4.44E-16	1.65E-15	AAEL005606-PA OS=Aedes aegypti GN=AAEL005606 PE=4 SV=1
CCG010726.2	1959	158	680	3.46565496	12.13457285	1.807923392	Up	0	0	AAEL006074-PA OS=Aedes aegypti GN=AAEL006074 PE=4 SV=1
CCG006223.1	1805	199	680	4.73738348	13.16987712	1.475079519	Up	0	0	AAEL011781-PA OS=Aedes aegypti GN=AAEL011781 PE=4 SV=1
CCG022868.1	3390	1761	680	22.3214462	7.012279707	-1.670475054	Down	6.26E-164	3.61E-163	AAEL012616-PA OS=Aedes aegypti GN=AAEL012616 PE=4 SV=1
CCG004056.1	1149	82	681	3.06659551	20.71939642	2.756272187	Up	0	0	-
CCG007692.1	1479	115	681	3.34112203	16.09640736	2.268334141	Up	6.13E-14	2.03E-13	AAEL012520-PA OS=Aedes aegypti GN=AAEL012520 PE=4 SV=1
CCG018303.1	1206	210	681	7.4822921	19.74012146	1.399578674	Up	0	0	AAEL002107-PA OS=Aedes aegypti GN=AAEL002107 PE=4 SV=1
CCG009518.1	1127	1144	682	43.6179027	21.15487556	-1.043930204	Down	1.42E-52	6.71E-52	AAEL007908-PA OS=Aedes aegypti GN=AAEL007908 PE=4 SV=1
CCG004536.1	2897	1943	682	28.8195355	8.22973585	-1.808129053	Down	6.14E-202	3.67E-201	AAEL013788-PA OS=Aedes aegypti GN=AAEL013788 PE=4 SV=1
CCG011670.1	2541	48	683	0.81170691	9.396498637	3.533092471	Up	5.35E-14	1.79E-13	AAEL004368-PA OS=Aedes aegypti GN=AAEL004368 PE=4 SV=1
CCG007349.1	1110	112	683	4.33568493	21.5103631	2.31070005	Up	0	0	Fanconi anemia group J protein OS=Danaus plexippus GN=KGM_13519 PE=4 SV=1
CCG004791.1	1176	175	683	6.39430575	20.30314884	1.66684386	Up	3.55E-13	9.58E-13	AAEL008743-PA OS=Aedes aegypti GN=AAEL008743 PE=4 SV=1

CCG015157.1	2261	216	683	4.10502551	10.56015172	1.36316747	Up	1.38E-13	4.17E-13	AGAP006748-PA OS=Anopheles gambiae GN=AGAP006748 PE=4 SV=2
CCG026234.1	2211	236	683	4.58654788	10.79896112	1.235411923	Up	4.60E-13	1.20E-12	AAEL005499-PA OS=Aedes aegypti GN=AAEL005499 PE=3 SV=1
CCG018888.1	40924	1219	683	1.27993614	0.583435222	-1.133427439	Down	1.09E-63	5.29E-63	AAEL011412-PA OS=Aedes aegypti GN=AAEL011412 PE=4 SV=1
CCG006192.1	2916	37	684	0.5452264	8.200089613	3.910712353	Up	5.17E-14	1.74E-13	WU-repeat protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011019 PE=4 SV=1
CCG009605.1	1623	177	684	4.68616329	14.7328782	1.652560169	Up	0	0	AAEL005584-PA OS=Aedes aegypti GN=AAEL005584 PE=4 SV=1
CCG000331.1	2522	1728	684	29.4415945	9.481150401	-1.634721783	Down	5.43E-156	3.11E-155	AAEL005977-PA OS=Aedes aegypti GN=AAEL005977 PE=4 SV=1
CCG012860.1	1902	65	685	1.46847148	12.59012597	3.099905569	Up	0	0	AAEL002921-PA OS=Aedes aegypti GN=AAEL002921 PE=4 SV=1
CCG007966.1	774	176	685	9.77089573	30.9385266	1.662841763	Up	0	0	AAEL002235-PA OS=Aedes aegypti GN=AAEL002235 PE=4 SV=1
CCG016668.1	1343	38	686	1.21582272	17.85657324	3.876450457	Up	4.26E-14	1.45E-13	AAEL000436-PA OS=Aedes aegypti GN=AAEL000436 PE=4 SV=1
CCG004869.1	1924	110	686	2.45668961	12.46433361	2.343018256	Up	0	0	AAEL007661-PA OS=Aedes aegypti GN=AAEL007661 PE=3 SV=1
CCG008875.1	849	157	686	7.94611111	28.24661704	1.829757221	Up	0	0	AAEL009220-PA OS=Aedes aegypti GN=AAEL009220 PE=4 SV=1
CCG017772.1	2568	206	686	3.44694912	9.338542783	1.437877443	Up	1.10E-13	3.44E-13	AAEL006226-PA OS=Aedes aegypti GN=AAEL006226 PE=4 SV=1
CCG009263.1	2307	214	686	3.9859225	10.39504892	1.382910984	Up	0	0	AAEL003205-PA OS=Aedes aegypti GN=AAEL003205 PE=4 SV=1
CCG003462.1	3384	231	686	2.9332177	7.086695587	1.272628929	Up	1.29E-13	3.97E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001735 PE=4 SV=1
CCG019227.1	2094	67	687	1.37486734	11.46911946	3.060390302	Up	0	0	AAEL001933-PA OS=Aedes aegypti GN=AAEL001933 PE=4 SV=1
CCG012968.1	1683	78	687	1.9914672	14.26995612	2.841077274	Up	0	0	AAEL013862-PA OS=Aedes aegypti GN=AAEL013862 PE=3 SV=1

CCG024274.1	1771	102	687	2.47482379	13.56088997	2.454054151	Up	0	0	AAEL008538-PA OS=Aedes aegypti GN=AAEL008538 PE=4 SV=1
CCG010443.1	2163	160	687	3.17852868	11.10325296	1.804551398	Up	4.24E-14	1.45E-13	AAEL007462-PA OS=Aedes aegypti GN=AAEL007462 PE=4 SV=1
CCG006893.1	1011	173	687	7.35288238	23.7550308	1.691851265	Up	3.60E-13	9.69E-13	AAEL014237-PA OS=Aedes aegypti GN=AAEL014237 PE=4 SV=1
CCG023914.1	1978	2973	687	64.5849449	12.14172707	-2.411224255	Down	0	0	AAEL003837-PA OS=Aedes aegypti GN=AAEL003837 PE=4 SV=1
CCG024611.1	1395	29	688	0.89327764	17.24107127	4.270596963	Up	6.66E-13	1.69E-12	AAEL009023-PA OS=Aedes aegypti GN=AAEL009023 PE=4 SV=1
CCG001902.1	1864	39	688	0.89904488	12.90305495	3.84317574	Up	7.33E-14	2.38E-13	AAEL012568-PA OS=Aedes aegypti GN=AAEL012568 PE=4 SV=1
CCG024552.1	1737	82	688	2.02850791	13.8464562	2.771025954	Up	0	0	AAEL003338-PA OS=Aedes aegypti GN=AAEL003338 PE=4 SV=1
CCG004151.1	1848	95	688	2.20894198	13.01476971	2.55872235	Up	0	0	AAEL006988-PA OS=Aedes aegypti GN=AAEL006988 PE=4 SV=1
CCG004175.1	810	105	688	5.57015078	29.69295607	2.414332441	Up	0	0	AAEL005304-PA OS=Aedes aegypti GN=AAEL005304 PE=4 SV=1
CCG022995.1	2205	158	688	3.07901046	10.90761652	1.82479721	Up	0	0	AAEL005067-PA OS=Aedes aegypti GN=AAEL005067 PE=4 SV=1
CCG003031.1	444	263	688	25.4527932	54.16958203	1.089658969	Up	1.40E-14	4.97E-14	AAEL008768-PA OS=Aedes aegypti GN=AAEL008768 PE=2 SV=1
CCG010650.1	446	1	689	0.0963447	54.00505089	9.130673377	Up	0.000587	0.00091723	Chromosome undetermined SCAF14565, whole genome shotgun sequence OS=Tetraodon nigroviridis GN=GSTENG00016696001 PE=3 SV=1
CCG018006.1	3864	77	689	0.85628094	6.233502251	2.863886836	Up	0	0	AAEL008021-PA OS=Aedes aegypti GN=AAEL008021 PE=4 SV=1
CCG013301.1	5388	99	689	0.78953299	4.47035128	2.501316756	Up	0	0	AAEL007856-PA OS=Aedes aegypti GN=AAEL007856 PE=4 SV=1
CCG004284.1	2817	122	689	1.86095407	8.550320446	2.199936039	Up	8.04E-14	2.58E-13	AAEL005719-PA OS=Aedes aegypti GN=AAEL005719 PE=4 SV=1

CCG021765.1	1999	155	689	3.33182034	12.04915092	1.854548971	Up	3.42E-13	9.26E-13	AAEL011420-PA OS=Aedes aegypti GN=AAEL011420 PE=4 SV=1
CCG022427.1	3390	29	690	0.3675877	7.115401467	4.27478476	Up	6.66E-13	1.69E-12	AAEL008126-PA OS=Aedes aegypti GN=GPRCIR PE=4 SV=2
CCG006646.1	1580	43	690	1.16942949	15.26658922	3.706501001	Up	8.46E-14	2.70E-13	AAEL013540-PA OS=Aedes aegypti GN=AAEL013540 PE=4 SV=1
CCG019326.1	264	63	690	10.2541412	91.36822338	3.155485832	Up	0	0	AAEL007125-PA OS=Aedes aegypti GN=AAEL007125 PE=4 SV=1
CCG017408.1	1074	232	690	9.28210282	22.45922809	1.27478476	Up	2.91E-13	8.05E-13	AAEL008700-PA OS=Aedes aegypti GN=AAEL008700 PE=4 SV=1
CCG011162.1	835	1126	690	57.9448158	28.88767781	-1.004225357	Down	1.16E-48	5.37E-48	AAEL006619-PA OS=Aedes aegypti GN=AAEL006619 PE=4 SV=1
CCG005498.1	774	14	691	0.77723034	31.20952099	5.327500182	Up	3.07E-08	6.01E-08	AAEL013692-PA OS=Aedes aegypti GN=PIWI3 PE=4 SV=1
CCG026182.1	1756	36	691	0.8809285	13.75636062	3.964930103	Up	6.13E-14	2.03E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013877 PE=4 SV=1
CCG027958.1	8953	188	691	0.90230203	2.69810893	1.580266253	Up	1.80E-13	5.25E-13	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1
CCG008226.1	1605	59	692	1.5795728	15.0723536	3.254298382	Up	0	0	AAEL003172-PA OS=Aedes aegypti GN=AAEL003172 PE=4 SV=1
CCG012478.1	1110	83	692	3.21305223	21.79380858	2.761902	Up	1.32E-13	4.01E-13	AAEL013006-PA OS=Aedes aegypti GN=AAEL013006 PE=4 SV=1
CCG013910.1	1524	119	692	3.35524831	15.87344326	2.242123668	Up	0	0	AAEL002784-PA OS=Aedes aegypti GN=AAEL002784 PE=4 SV=1
CCG025942.1	1942	62	693	1.37184529	12.47481246	3.184828436	Up	0	0	AAEL009281-PA OS=Aedes aegypti GN=AAEL009281 PE=4 SV=1
CCG002709.1	1757	76	693	1.85867947	13.7883243	2.891097233	Up	0	0	AAEL007573-PA OS=Aedes aegypti GN=AAEL007573 PE=4 SV=1
CCG005254.1	1005	143	693	6.11410154	24.10555801	1.979153409	Up	0	0	AAEL012749-PA OS=Aedes aegypti GN=AAEL012749 PE=4 SV=1
CCG011884.2	1176	148	693	5.40775572	20.6004131	1.92957138	Up	0	0	AAEL000204-PA OS=Aedes aegypti GN=AAEL000204 PE=4 SV=1
CCG002435.1	3561	276	693	3.33042593	6.803169279	1.030500289	Up	0	0	AAEL011162-PA OS=Aedes aegypti GN=AAEL011162 PE=4 SV=1

CCG001503.1	1905	59	694	1.33082118	12.73545621	3.258462007	Up	0	0	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006823 PE=4 SV=1
CCG026312.2	4572	157	694	1.47555738	5.306440087	1.846484308	Up	0	0	AAEL007215-PA OS=Aedes aegypti GN=AAEL007215 PE=4 SV=1
CCG013155.2	1242	201	694	6.95403918	19.53385192	1.490053365	Up	0	0	AAEL004449-PA OS=Aedes aegypti GN=AAEL004449 PE=4 SV=1
CCG016305.1	831	220	694	11.3758624	29.19499889	1.359745343	Up	4.61E-13	1.20E-12	AAEL006001-PA OS=Aedes aegypti GN=AAEL006001 PE=4 SV=1
CCG028399.1	2207	226	694	4.40016313	10.99277031	1.320926094	Up	2.04E-14	7.16E-14	AAEL006449-PA OS=Aedes aegypti GN=AAEL006449 PE=4 SV=1
CCG027466.1	2426	243	694	4.30405833	10.00043037	1.216292553	Up	1.53E-14	5.41E-14	AAEL011370-PA OS=Aedes aegypti GN=AAEL011370 PE=4 SV=1
CCG000807.1	3967	277	694	3.00040748	6.115715674	1.02736289	Up	0	0	AAEL012583-PA (Fragment) OS=Aedes aegypti GN=AAEL012583 PE=4 SV=1
CCG019780.1	2487	2610	694	45.0948964	9.755144383	-2.208729035	Down	0	0	AAEL003129-PA OS=Aedes aegypti GN=AAEL003129 PE=4 SV=1
CCG019334.1	411	5206	694	544.283305	59.02930433	-3.20485454	Down	0	0	AAEL013613-PA OS=Aedes aegypti GN=AAEL013613 PE=4 SV=1
CCG013418.2	1578	145	695	3.94842302	15.39670618	1.963273281	Up	0	0	AAEL007208-PA OS=Aedes aegypti GN=AAEL007208 PE=4 SV=1
CCG014737.1	792	201	695	10.9051978	30.67677065	1.49213068	Up	0	0	AAEL006845-PA OS=Aedes aegypti GN=AAEL006845 PE=4 SV=1
CCG027485.1	986	241	695	10.5027445	24.64097602	1.230293035	Up	2.37E-13	6.72E-13	AAEL000184-PA OS=Aedes aegypti GN=AAEL000184 PE=4 SV=1
CCG026708.2	2280	79	696	1.48886361	10.67147396	2.841475951	Up	0	0	AAEL013110-PA OS=Aedes aegypti GN=AAEL013110 PE=4 SV=1
CCG020432.1	1041	134	696	5.53116661	23.37268072	2.079167509	Up	0	0	RNA-interacting protein alpha OS=Culex quinquefasciatus GN=CpipJ_CPIJ008848 PE=4 SV=1
CCG011787.1	906	178	696	8.44217744	26.85536494	1.669523269	Up	1.15E-13	3.57E-13	AAEL012917-PA (Fragment) OS=Aedes aegypti GN=AAEL012917 PE=4 SV=1

CCG008624.1	2244	240	696	4.59569354	10.84267408	1.238366104	Up	0	0	AAEL007251-PA OS=Redes SV-1 aegypti GN=AAEL007251 PE=4
CCG020281.1	561	7513	696	575.457426	43.37069632	-3.729916687	Down	0	0	AAEL008490-PA OS=Redes SV-1 aegypti GN=AAEL008490 PE=4
CCG004951.1	465	4	697	0.36963213	52.39982561	7.14732805	Up	2.95E-05	5.03E-05	AAEL002047-PA OS=Redes SV-1 aegypti GN=AAEL002047 PE=2
CCG003067.1	1584	104	697	2.8212452	15.38252456	2.446888332	Up	0	0	-
CCG014887.1	558	115	697	8.85576968	43.66652134	2.301837999	Up	6.13E-14	2.03E-13	AAEL013176-PA OS=Redes SV-1 aegypti GN=AAEL013176 PE=4
CCG015132.1	2100	130	697	2.66003119	11.60281853	2.124960237	Up	0	0	AAEL011415-PA OS=Redes SV-1 aegypti GN=AAEL011415 PE=4
CCG027193.1	1905	145	697	3.27066221	12.79050861	1.96741896	Up	0	0	AAEL003679-PA OS=Redes SV-1 aegypti GN=AAEL003679 PE=4
CCG023761.1	439	3369	697	329.760902	55.50323214	-2.570776663	Down	0	0	-
CCG013230.1	481	0	698	0.001	50.7294744	15.63053659	Up	4.44E-16	1.63E-15	-
CCG018501.1	1558	0	698	0.001	15.661667	13.93495016	Up	4.44E-16	1.64E-15	AAEL005916-PA OS=Redes SV-1 aegypti GN=AAEL005916 PE=4
CCG006500.1	1992	97	698	2.09240174	12.24943634	2.549483588	Up	3.62E-14	1.25E-13	AAEL012392-PA OS=Redes SV-1 aegypti GN=AAEL012392 PE=4
CCG019959.1	1797	130	698	3.10855064	13.57867401	2.127028617	Up	0	0	AAEL008897-PA OS=Redes SV-1 aegypti GN=AAEL008897 PE=4
CCG020844.1	612	137	698	9.6190419	39.87071436	2.051364347	Up	2.90E-13	8.03E-13	AAEL014324-PA OS=Redes SV-1 aegypti GN=AAEL014324 PE=4
CCG007731.1	888	183	698	8.85524936	27.4784653	1.633696592	Up	3.05E-13	8.41E-13	AAEL006956-PA OS=Redes SV-1 aegypti GN=AAEL006956 PE=4
CCG021791.2	710	208	698	12.5883166	34.36743266	1.448956712	Up	3.93E-13	1.05E-12	AAEL006628-PA OS=Redes SV-1 aegypti GN=AAEL006628 PE=4
CCG017133.1	1329	2	699	0.06466476	18.38663316	8.151461849	Up	9.36E-05	0.00015344	AAEL009987-PA OS=Redes SV-1 aegypti GN=AAEL009987 PE=4
CCG000229.1	741	67	699	3.88525266	32.97683598	3.085372659	Up	0	0	AAEL003360-PA OS=Redes SV-1 aegypti GN=AAEL003360 PE=4
CCG018116.1	2134	87	699	1.75181205	11.45071952	2.708518353	Up	0	0	AAEL003273-PA OS=Redes SV-1 aegypti GN=AAEL003273 PE=4
CCG006585.1	981	1203	699	52.6937724	24.90910853	-1.080959078	Down	2.43E-58	1.16E-57	AAEL006065-PA OS=Redes SV-1 aegypti GN=AAEL006065 PE=3

CCG012429.1	1027	98	700	4.10032521	23.82745252	2.538814472	Up	0	0	Dihydrodipicolate reductase OS=Culex quinquefasciatus GN=CpipJ_CPIJ002390 PE=4 SV=1
CCG026004.1	990	116	700	5.03483759	24.71797348	2.295543321	Up	1.16E-13	3.58E-13	AAEL012874-PA OS=Aedes aegypti GN=AAEL012874 PE=4 SV=1
CCG019330.1	2688	155	700	2.47779348	9.103717909	1.87739991	Up	3.42E-13	9.27E-13	AAEL007130-PA OS=Aedes aegypti GN=AAEL007130 PE=3 SV=1
CCG003110.1	1311	210	700	6.88302385	18.66574656	1.439278798	Up	0	0	AAEL002102-PA OS=Aedes aegypti GN=AAEL002102 PE=4 SV=1
CCG008297.1	3042	96	701	1.35604685	8.055802767	2.570621337	Up	0	0	Rhopper OS=Culex quinquefasciatus GN=CpipJ_CPIJ004355 PE=4 SV=1
CCG008972.1	2412	189	702	3.36703144	10.17442384	1.595398	Up	2.34E-13	6.64E-13	DNA topoisomerase OS=Aedes aegypti GN=AAEL011603 PE=3 SV=1
CCG016992.2	12986	2545	702	8.4212209	1.889782096	-2.155809517	Down	0	0	AAEL010187-PA (Fragment) OS=Aedes aegypti GN=AAEL010187 PE=4 SV=1
CCG002130.1	1401	88	703	2.69902687	17.54151932	2.700262464	Up	2.93E-14	1.02E-13	AAEL013540-PA OS=Aedes aegypti GN=AAEL013540 PE=4 SV=1
CCG012542.2	1710	93	703	2.33695048	14.37173601	2.620535272	Up	4.04E-14	1.39E-13	AAEL005571-PA OS=Aedes aegypti GN=AAEL005571 PE=4 SV=1
CCG018045.1	627	112	703	7.67561448	39.19564365	2.352339161	Up	0	0	Subolesin-like protein OS=Aedes albopictus PE=2 SV=1
CCG026667.1	2225	135	703	2.60715244	11.0452443	2.082878486	Up	1.90E-13	5.48E-13	AAEL002063-PA OS=Aedes aegypti GN=AAEL002063 PE=4 SV=1
CCG004838.1	1299	142	703	4.69723042	18.91891345	2.009946963	Up	2.62E-13	7.34E-13	AAEL010107-PA OS=Aedes aegypti GN=AAEL010107 PE=4 SV=1
CCG019575.1	2034	168	703	3.54912262	12.08243293	1.76737666	Up	1.45E-13	4.33E-13	AAEL001765-PA OS=Aedes aegypti GN=AAEL001765 PE=4 SV=1
CCG013402.1	1719	260	703	6.49920361	14.29649132	1.13732627	Up	3.22E-13	8.81E-13	AGAP010551-PA (Fragment) OS=Anopheles gambiae GN=AGAP010551 PE=4 SV=4
CCG011613.2	1514	273	703	7.7481754	16.23227779	1.066936942	Up	1.53E-14	5.41E-14	AAEL012586-PA OS=Aedes aegypti GN=AAEL012586 PE=4 SV=1

CCG017294.1	941	1287	703	58.7694457	26.11654471	-1.170102255	Down	1.38E-70	6.85E-70	AAEL012023-PA OS=Aedes aegypti GN=AAEL012023 PE=4 SV=1
CCG006299.1	2508	47	704	0.8052542	9.81284962	3.607155971	Up	0	0	AAEL005281-PA OS=Aedes aegypti GN=AAEL005281 PE=4 SV=1
CCG005103.1	639	127	704	8.5401507	38.51428302	2.173060136	Up	1.57E-13	4.66E-13	AAEL006908-PA OS=Aedes aegypti GN=AAEL006908 PE=4 SV=1
CCG014291.1	1440	173	704	5.16233617	17.09071309	1.727116595	Up	3.60E-13	9.69E-13	AAEL012016-PA OS=Aedes aegypti GN=AAEL012016 PE=4 SV=1
CCG020803.1	1011	200	704	8.50044206	24.34285544	1.517888633	Up	0	0	AAEL001158-PB OS=Aedes aegypti GN=AAEL001158 PE=3 SV=1
CCG027344.1	521	2324	704	191.673058	47.23728762	-2.020649531	Down	2.82E-280	1.76E-279	-
CCG004234.1	936	27	705	1.23951158	26.33075334	4.408905149	Up	2.28E-12	5.41E-12	AAEL008124-PA OS=Aedes aegypti GN=AAEL008124 PE=4 SV=1
CCG022301.1	1291	33	705	1.09837432	19.09030606	4.119398532	Up	1.31E-13	4.00E-13	AAEL011762-PA OS=Aedes aegypti GN=AAEL011762 PE=4 SV=1
CCG017013.1	1305	52	705	1.71220398	18.88550584	3.463352933	Up	0	0	Polyadenylate-binding protein-like protein OS=Aedes aegypti PE=2 SV=1
CCG000243.1	894	168	705	8.07484946	27.5677686	1.771475228	Up	1.45E-13	4.33E-13	AAEL007179-PA OS=Aedes aegypti GN=AAEL007179 PE=4 SV=1
CCG006321.1	1752	176	705	4.31659434	14.0671148	1.704361032	Up	0	0	AAEL000426-PA OS=Aedes aegypti GN=AAEL000426 PE=4 SV=1
CCG017560.1	651	1342	705	88.5796987	37.85804167	-1.226376305	Down	4.11E-79	2.08E-78	AAEL007410-PA OS=Aedes aegypti GN=AAEL007410 PE=4 SV=1
CCG000228.1	543	1485	705	117.513915	45.38781791	-1.372454565	Down	8.35E-104	4.45E-103	AAEL006652-PA OS=Aedes aegypti GN=AAEL006652 PE=4 SV=1
CCG004142.1	1323	122	706	3.96243962	18.65498367	2.23510024	Up	8.04E-14	2.58E-13	AAEL007560-PA OS=Aedes aegypti GN=AAEL007560 PE=4 SV=1
CCG015988.1	1572	160	706	4.37350988	15.70009122	1.843909482	Up	4.24E-14	1.44E-13	Zinc finger protein 189 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000772 PE=4 SV=1
CCG028447.1	1278	93	707	3.12690557	19.33920319	2.628720798	Up	4.04E-14	1.38E-13	AAEL014536-PA OS=Aedes aegypti GN=AAEL014536 PE=4 SV=1
CCG022160.1	2333	113	707	2.08126018	10.59387127	2.347700646	Up	1.42E-13	4.26E-13	AAEL013552-PA OS=Aedes aegypti GN=AAEL013552 PE=4 SV=1

CCG005637.1	1410	3091	707	94.1981913	17.52872459	-2.42597833	Down	0	0	AAEL010075-PA OS=Aedes aegypti GN=AAEL010075 PE=3 SV=1
CCG028283.1	591	34	708	2.47203211	41.87895085	4.082455913	Up	8.82E-14	2.80E-13	AAEL010109-PA OS=Aedes aegypti GN=AAEL010109 PE=4 SV=1
CCG014839.1	1052	85	708	3.4718892	23.52705319	2.760527818	Up	0	0	AAEL011700-PA OS=Aedes aegypti GN=AAEL011700 PE=4 SV=1
CCG022754.1	9760	2838	708	12.4946831	2.535907782	-2.30074012	Down	0	0	AAEL008478-PA OS=Aedes aegypti GN=AAEL008478 PE=4 SV=1
CCG000917.1	936	33	709	1.51495859	26.48014768	4.127560902	Up	1.31E-13	4.01E-13	AAEL015338-PA OS=Aedes aegypti GN=AAEL015338 PE=4 SV=1
CCG028467.1	1059	60	709	2.43454587	23.40454979	3.265064425	Up	0	0	AAEL010747-PA OS=Aedes aegypti GN=AAEL010747 PE=4 SV=1
CCG006687.1	1395	69	709	2.12538472	17.7673249	3.063430564	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016357 PE=4 SV=1
CCG022437.1	1886	95	709	2.1644352	13.14179122	2.602099413	Up	0	0	Zinc finger protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010644 PE=4 SV=1
CCG001106.1	1587	85	710	2.30146657	15.63980876	2.764597482	Up	0	0	AAEL007368-PA OS=Aedes aegypti GN=AAEL007368 PE=4 SV=1
CCG001102.1	1084	116	710	4.59823728	22.8970263	2.316007423	Up	1.16E-13	3.59E-13	AAEL006144-PA OS=Aedes aegypti GN=AAEL006144 PE=2 SV=1
CCG013345.1	930	229	710	10.5807196	26.68857689	1.33478463	Up	2.46E-13	6.93E-13	AAEL004797-PA OS=Aedes aegypti GN=AAEL004797 PE=4 SV=1
CCG025310.4	1686	40	711	1.01944803	14.74219145	3.854090859	Up	6.86E-14	2.24E-13	AAEL006681-PA OS=Aedes aegypti GN=AAEL006681 PE=4 SV=1
CCG005638.1	1487	203	711	5.86607675	16.71508728	1.510683036	Up	9.28E-14	2.93E-13	AAEL009079-PA OS=Aedes aegypti GN=AAEL009079 PE=4 SV=1
CCG009091.1	1011	12	712	0.51002652	24.6194788	5.593084134	Up	1.30E-07	2.49E-07	AAEL004593-PA OS=Aedes aegypti GN=AAEL004593 PE=4 SV=1
CCG000493.1	1777	63	712	1.52340646	14.00691787	3.200766711	Up	0	0	AAEL001580-PA OS=Aedes aegypti GN=AAEL001580 PE=4 SV=1
CCG002336.1	2292	95	712	1.78103176	10.85963921	2.608191026	Up	0	0	Zinc finger protein 558 OS=Culex quinquefasciatus GN=CpipJ_CPIJ018144 PE=4 SV=1

CCG000653.1	604	122	712	8.67931726	41.20909447	2.247309297	Up	8.04E-14	2.58E-13	AAEL004343-PA OS=Aedes aegypti GN=AAEL004345 PE=3 SV=1
CCG012619.1	1086	3006	712	118.938326	22.91923855	-2.375582659	Down	0	0	AAEL007542-PA OS=Aedes aegypti GN=AAEL007542 PE=3 SV=1
CCG003163.1	1158	31	713	1.15031241	21.52439667	4.22587516	Up	2.40E-13	6.78E-13	AAEL004316-PB OS=Aedes aegypti GN=AAEL004318 PE=4 SV=1
CCG009248.1	793	151	713	8.18213105	31.43159059	1.941666731	Up	0	0	GK19160 OS=Drosophila willistoni GN=GK19160 PE=4 SV=1
CCG012081.1	3346	260	713	3.33895128	7.449268183	1.157703657	Up	3.22E-13	8.82E-13	AAEL012449-PA OS=Aedes aegypti GN=AAEL012449 PE=4 SV=1
CCG003498.1	2022	74	714	1.57258178	12.34431732	2.972640102	Up	0	0	AAEL007642-PA OS=Aedes aegypti GN=TUBE PE=4 SV=1
CCG013120.1	943	128	714	5.83258328	26.46893915	2.182093468	Up	4.15E-14	1.42E-13	AAEL012624-PA OS=Aedes aegypti GN=AAEL012624 PE=4 SV=1
CCG001114.1	1002	168	714	7.2045064	24.91038884	1.789776045	Up	1.45E-13	4.33E-13	AAEL000883-PA OS=Aedes aegypti GN=AAEL000883 PE=4 SV=1
CCG012595.1	2609	226	714	3.72217709	9.566964207	1.361914505	Up	2.04E-14	7.16E-14	Uncharacterized protein OS=Anopheles darlingi GN=AND_10270 PE=4 SV=1
CCG020007.1	1630	256	714	6.7486209	15.31301203	1.182093468	Up	0	0	AAEL014283-PA OS=Aedes aegypti GN=AAEL014283 PE=4 SV=1
CCG023188.1	743	8	715	0.46266201	33.64087199	6.184112635	Up	2.19E-06	3.96E-06	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013878 PE=4 SV=1
CCG016251.1	858	115	715	5.75934671	29.13189731	2.338622585	Up	6.13E-14	2.03E-13	Putative uncharacterized protein OS=Amblyomma maculatum PE=2 SV=1
CCG008040.1	3658	170	715	1.99695322	6.833014733	1.774721699	Up	3.53E-13	9.53E-13	AAEL001370-PA OS=Aedes aegypti GN=AAEL001370 PE=3 SV=1
CCG017222.1	1976	204	715	4.43614669	12.64937646	1.511687294	Up	6.59E-14	2.17E-13	AAEL009374-PA OS=Aedes aegypti GN=AAEL009374 PE=4 SV=1
CCG019905.1	1782	86	716	2.07373579	14.04608651	2.759864226	Up	0	0	Small nuclear ribonucleoprotein-associated protein OS=Aedes aegypti GN=AAEL002377 PE=2 SV=1

CCG012793.1	1515	125	716	3.54535764	16.52153543	2.220344696	Up	2.35E-14	8.23E-14	AAEL011862-PA OS=Aedes aegypti GN=AAEL011862 PE=4 SV=1
CCG012302.1	2648	82	717	1.33063378	9.465666331	2.830590508	Up	0	0	AAEL008463-PA OS=Aedes aegypti GN=AAEL008463 PE=3 SV=1
CCG020325.1	1191	230	717	8.29810156	21.04541095	1.342652462	Up	3.85E-13	1.03E-12	AAEL012209-PA OS=Aedes aegypti GN=AAEL012209 PE=4 SV=1
CCG003277.1	1360	179	718	5.65557536	18.45591377	1.70633746	Up	0	0	AAEL007895-PA OS=Aedes aegypti GN=AAEL007895 PE=4 SV=1
CCG002929.1	2268	181	718	3.42924249	11.06703824	1.690307351	Up	1.53E-13	4.55E-13	AAEL000315-PA OS=Aedes aegypti GN=AAEL000315 PE=4 SV=1
CCG022396.1	1281	214	718	7.17839438	19.59410049	1.448686251	Up	0	0	AAEL008005-PA OS=Aedes aegypti GN=AAEL008005 PE=4 SV=1
CCG023977.1	1506	68	719	1.9402005	16.6899077	3.104698323	Up	0	0	AAEL000260-PA OS=Aedes aegypti GN=AAEL000260 PE=4 SV=1
CCG025401.1	2721	143	719	2.25824037	9.237413083	2.032289827	Up	0	0	AAEL005837-PA OS=Aedes aegypti GN=AAEL005837 PE=4 SV=1
CCG015645.1	2733	180	719	2.83005936	9.19685364	1.700308068	Up	3.32E-13	9.05E-13	AAEL009261-PA OS=Aedes aegypti GN=AAEL009261 PE=4 SV=1
CCG006750.1	2514	232	719	3.96538521	9.998011535	1.334180169	Up	2.91E-13	8.05E-13	AGAP008245-PA (Fragment) OS=Anopheles gambiae GN=AGAP008245 PE=4 SV=3
CCG002163.1	1644	1391	719	36.3569956	15.28893005	-1.24974554	Down	2.30E-84	1.18E-83	AAEL005666-PA OS=Aedes aegypti GN=AAEL005666 PE=4 SV=1
CCG014702.1	2328	5146	719	94.9837862	10.79682174	-3.137074577	Down	0	0	AAEL002971-PA OS=Aedes aegypti GN=AAEL002971 PE=4 SV=1
CCG020005.1	612	20	720	1.40423969	41.12738444	4.872238205	Up	3.69E-10	7.72E-10	AAEL014287-PA OS=Aedes aegypti GN=AAEL014287 PE=3 SV=1
CCG023857.1	2673	93	720	1.49501882	9.416370848	2.655007489	Up	4.04E-14	1.39E-13	AAEL008550-PA OS=Aedes aegypti GN=AAEL008550 PE=4 SV=1
CCG023181.1	1128	189	720	7.19971617	22.31379368	1.631923876	Up	2.34E-13	6.64E-13	Rab-protein OS=Aedes aegypti PE=2 SV=1
CCG007047.1	636	190	720	12.8368704	39.57540767	1.624310692	Up	2.69E-13	7.52E-13	AAEL001615-PA OS=Aedes aegypti GN=AAEL001615 PE=4 SV=1
CCG025702.1	2403	275	720	4.91746859	10.47439004	1.090878492	Up	4.31E-13	1.14E-12	AAEL012969-PA OS=Aedes aegypti GN=AAEL012969 PE=4 SV=1

CCG015517.1	2561	263	721	4.4127451	9.841826456	1.157249664	Up	1.40E-14	4.97E-14	AAEL005402-PA OS=Aedes aegypti GN=AAEL005402 PE=4 SV=1
CCG008026.1	851	14	722	0.70690515	29.65907853	5.390813309	Up	3.07E-08	6.01E-08	AAEL005000-PA OS=Aedes aegypti GN=AAEL005000 PE=4 SV=1
CCG001644.1	1560	21	722	0.57843874	16.17940758	4.805850808	Up	1.76E-10	3.72E-10	AAEL000739-PA OS=Aedes aegypti GN=AAEL000739 PE=4 SV=1
CCG026526.1	1131	52	722	1.97561998	22.31642425	3.497728513	Up	0	0	AAEL011659-PA OS=Aedes aegypti GN=AAEL011659 PE=4 SV=1
CCG017580.1	582	72	722	5.31584346	43.36748424	3.028243229	Up	0	0	AAEL002103-PA OS=Aedes aegypti GN=AAEL002103 PE=4 SV=1
CCG007398.1	1599	123	722	3.3053642	15.78478789	2.255653725	Up	0	0	AAEL004057-PA OS=Aedes aegypti GN=AAEL004057 PE=4 SV=1
CCG028378.1	1078	218	722	8.68961238	23.41361394	1.429983906	Up	0	0	-
CCG004481.1	816	102	723	5.37121683	30.9740614	2.527739699	Up	0	0	AAEL002103-PA OS=Aedes aegypti GN=AAEL002103 PE=4 SV=1
CCG020809.1	975	190	723	8.37358931	25.92290678	1.630309432	Up	2.69E-13	7.52E-13	AAEL001154-PA OS=Aedes aegypti GN=AAEL001154 PE=4 SV=1
CCG018820.3	1410	1184	723	36.0823871	17.92541426	-1.009288325	Down	1.72E-51	8.05E-51	Tubulin beta-1 chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ003635 PE=3 SV=1
CCG015242.2	6256	2373	723	16.2991017	4.040094966	-2.012331344	Down	1.38E-284	8.62E-284	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007959 PE=4 SV=1
CCG018480.1	785	171	724	9.36028614	32.24177374	1.784306576	Up	3.42E-13	9.25E-13	Ribulose-phosphate 3-epimerase OS=Aedes aegypti PE=2 SV=1
CCG016608.2	2070	276	724	5.72929795	12.22695284	1.093634634	Up	0	0	Ceramide kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ003342 PE=4 SV=1
CCG023297.1	1968	1850	724	40.3932973	12.86066686	-1.651150465	Down	2.82E-169	1.64E-168	Discs overgrown protein kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ006381 PE=4 SV=1
CCG019434.1	974	21	725	0.92645218	26.02130458	4.811832966	Up	1.76E-10	3.73E-10	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005419 PE=4 SV=1

CCG011141.1	1218	126	725	4.44514496	20.80849808	2.226870465	Up	1.34E-13	4.06E-13	AAEL000956-PA OS=Reedes aegypti GN=AAEL000956 PE=4 SV=1
CCG028131.1	984	181	725	7.90398574	25.75686043	1.704304502	Up	1.53E-13	4.55E-13	AAEL007370-PA OS=Reedes aegypti GN=AAEL007370 PE=3 SV=1
CCG025352.1	2664	128	726	2.06461187	9.526917769	2.206138942	Up	4.15E-14	1.42E-13	AAEL011811-PA OS=Reedes aegypti GN=AAEL011811 PE=3 SV=1
CCG027257.1	942	177	726	8.07393103	26.94236617	1.738533392	Up	0	0	AAEL001709-PA OS=Reedes aegypti GN=AAEL001709 PE=4 SV=1
CCG008489.1	714	255	726	15.3463338	35.54581084	1.211785505	Up	0	0	AAEL004445-PA OS=Reedes aegypti GN=AAEL004445 PE=4 SV=1
CCG023634.1	1761	119	727	2.90369019	14.43195185	2.313306994	Up	0	0	AAEL007943-PA OS=Reedes aegypti GN=AAEL007943 PE=4 SV=1
CCG009827.1	1530	135	727	3.79144717	16.6108936	2.131309161	Up	1.90E-13	5.49E-13	AAEL014400-PA OS=Reedes aegypti GN=AAEL014400 PE=4 SV=1
CCG009174.1	1194	1779	727	64.0227453	21.28531592	-1.588726038	Down	4.68E-154	2.67E-153	AAEL005319-PA OS=Reedes aegypti GN=AAEL005319 PE=4 SV=1
CCG025294.1	2025	21	728	0.44561206	12.56771629	4.817790421	Up	1.76E-10	3.72E-10	AAEL006549-PA OS=Reedes aegypti GN=AAEL006549 PE=4 SV=1
CCG003972.1	1509	72	728	2.05024579	16.86522564	3.040182843	Up	0	0	AAEL010447-PA OS=Reedes aegypti GN=AAEL010447 PE=4 SV=1
CCG021179.1	1257	128	728	4.37559748	20.24632099	2.210107844	Up	4.15E-14	1.42E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_22964 PE=4 SV=1
CCG012566.1	1077	204	729	8.13911408	23.66256617	1.539662866	Up	6.59E-14	2.17E-13	-
CCG019486.1	2100	253	729	5.17682993	12.13551608	1.229094633	Up	3.13E-13	8.61E-13	AAEL006075-PA OS=Reedes aegypti GN=AAEL006075 PE=4 SV=1
CCG000895.1	846	270	729	13.7137451	30.12362147	1.135272611	Up	2.59E-13	7.26E-13	AAEL015284-PA OS=Reedes aegypti GN=AAEL015284 PE=4 SV=1
CCG001467.1	3579	26	730	0.31215789	7.130355419	4.513626139	Up	4.60E-12	1.05E-11	AAEL012489-PA OS=Reedes aegypti GN=AAEL012489 PE=4 SV=1
CCG018475.1	996	43	730	1.85511906	25.62203016	3.787801103	Up	8.46E-14	2.70E-13	AAEL007167-PA OS=Reedes aegypti GN=AAEL007167 PE=4 SV=1
CCG017812.1	2524	145	730	2.46854656	10.11075358	2.034156768	Up	0	0	AAEL012202-PA OS=Reedes aegypti GN=AAEL012202 PE=4 SV=1

CCG021001.1	1269	6	731	0.20316659	20.1375101	6.631078299	Up	8.38E-06	1.47E-05	AAEL013830-PA OS=Aedes aegypti GN=AAEL013830 PE=4 SV=1
CCG005262.1	3045	110	731	1.55227284	8.392282535	2.434681086	Up	0	0	AAEL004441-PA OS=Aedes aegypti GN=AAEL004441 PE=4 SV=1
CCG020063.1	2037	216	731	4.55643725	12.54516461	1.461153298	Up	1.38E-13	4.17E-13	AAEL014149-PA OS=Aedes aegypti GN=AAEL014149 PE=4 SV=1
CCG001812.1	552	263	731	20.4728989	46.29438464	1.17712181	Up	1.40E-14	4.97E-14	AAEL005816-PA OS=Aedes aegypti GN=AAEL005816 PE=4 SV=1
CCG017723.1	621	36	732	2.49099911	41.20685764	4.048088041	Up	6.13E-14	2.03E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014666 PE=4 SV=1
CCG025635.1	2055	99	732	2.07007481	12.45229129	2.588656422	Up	0	0	AAEL010663-PA OS=Aedes aegypti GN=AAEL010663 PE=4 SV=1
CCG001050.1	1468	101	732	2.95636457	17.43151131	2.559801559	Up	0	0	AAEL002772-PA OS=Aedes aegypti GN=AAEL002772 PE=4 SV=1
CCG021181.2	1751	101	732	2.47855123	14.6141968	2.559801559	Up	0	0	AAEL002366-PA OS=Aedes aegypti GN=AAEL002366 PE=4 SV=1
CCG024570.2	2115	214	732	4.34776511	12.0990348	1.476546056	Up	0	0	AAEL002782-PA (Fragment) OS=Aedes aegypti GN=AAEL002782 PE=4 SV=1
CCG007597.1	2264	85	733	1.613263	11.31820533	2.810591656	Up	0	0	AAEL010650-PA OS=Aedes aegypti GN=AAEL010650 PE=3 SV=1
CCG013999.1	1551	152	733	4.2110894	16.521223	1.972055079	Up	8.39E-14	2.69E-13	AAEL003770-PA OS=Aedes aegypti GN=AAEL003770 PE=3 SV=1
CCG008897.1	982	171	733	7.48250979	26.09411087	1.802130077	Up	3.42E-13	9.27E-13	AAEL000844-PA OS=Aedes aegypti GN=AAEL000844 PE=4 SV=1
CCG025278.1	1122	197	733	7.5445969	22.83816121	1.597930772	Up	0	0	Structural maintenance of chromosomes protein OS=Aedes aegypti GN=AAEL001655 PE=3 SV=1
CCG002476.1	1140	248	733	9.34780192	22.47755866	1.265786282	Up	5.86E-14	1.95E-13	AAEL009767-PA OS=Aedes aegypti GN=AAEL009767 PE=4 SV=1
CCG024055.1	2910	44	734	0.6497142	8.817654691	3.762517838	Up	3.91E-14	1.34E-13	AAEL013016-PA OS=Aedes aegypti GN=AAEL013016 PE=4 SV=1

CCG025705.1	2286	128	734	2.40600439	11.22457356	2.221949457	Up	4.15E-14	1.42E-13	AAEL013594-PA OS=Aedes aegypti GN=AAEL013594 PE=4 SV=1
CCG012623.1	1460	145	734	4.26754214	17.57491449	2.042040367	Up	0	0	AAEL007541-PA OS=Aedes aegypti GN=AAEL007541 PE=4 SV=1
CCG017559.1	2967	155	734	2.24479571	8.648255865	1.945825051	Up	3.42E-13	9.26E-13	AAEL007409-PA (Fragment) OS=Aedes aegypti GN=AAEL007409 PE=4 SV=1
CCG016365.1	3429	1577	734	19.7618173	7.483049038	-1.401017488	Down	1.35E-113	7.32E-113	AAEL004694-PA OS=Aedes aegypti GN=AAEL004694 PE=4 SV=1
CCG000524.1	2481	40	735	0.69278089	10.35644233	3.901985549	Up	6.86E-14	2.24E-13	AAEL010280-PA OS=Aedes aegypti GN=AAEL010280 PE=4 SV=1
CCG025761.1	2796	84	735	1.29093623	9.189675761	2.831596221	Up	0	0	AAEL010497-PA OS=Aedes aegypti GN=AAEL010497 PE=4 SV=1
CCG000357.1	561	115	735	8.80841262	45.80095085	2.378423593	Up	6.13E-14	2.03E-13	AGAP003741-PA OS=Anopheles gambiae GN=AgaP_AGAP003741 PE=4 SV=3
CCG027541.1	1113	119	735	4.59424835	23.08565447	2.32909588	Up	0	0	AAEL012913-PA OS=Aedes aegypti GN=AAEL012913 PE=4 SV=1
CCG028619.1	1011	217	735	9.22297963	25.41477095	1.462362411	Up	1.47E-13	4.37E-13	AAEL014237-PA OS=Aedes aegypti GN=AAEL014237 PE=4 SV=1
CCG019691.1	1645	236	735	6.16465494	15.61965558	1.341270594	Up	4.60E-13	1.20E-12	AAEL005557-PA OS=Aedes aegypti GN=AAEL005557 PE=4 SV=1
CCG022072.1	1785	256	735	6.1626062	14.39458455	1.223913644	Up	0	0	AAEL004916-PA OS=Aedes aegypti GN=AAEL004916 PE=4 SV=1
CCG023420.1	5223	3609	735	29.6913215	4.919458822	-2.593469784	Down	0	0	AAEL002972-PA OS=Aedes aegypti GN=AAEL002972 PE=3 SV=1
CCG004886.1	759	97	736	5.49152076	33.89893505	2.625962318	Up	3.62E-14	1.25E-13	AAEL015658-PA OS=Aedes aegypti GN=AAEL015658 PE=4 SV=1
CCG021369.1	717	134	736	8.03060591	35.88464673	2.159785969	Up	0	0	AAEL004437-PA OS=Aedes aegypti GN=AAEL004437 PE=4 SV=1
CCG004942.1	2221	201	736	3.88875131	11.58455277	1.574823469	Up	0	0	AAEL010761-PA OS=Aedes aegypti GN=AAEL010761 PE=4 SV=1

CCG021223.1	31651	1697	736	2.30386527	0.812906123	-1.50289569	Down	1.43E-135	8.01E-135	AGAP012613-PA OS=Anopheles gambiae str. PEST GN=AgaP_AGAP012613 PE=4 SV=1
CCG001846.1	951	125	737	5.64796722	27.09174551	2.262049728	Up	2.35E-14	8.22E-14	AAEL007258-PA OS=Aedes aegypti GN=AAEL007258 PE=4 SV=1
CCG007363.1	1399	200	737	6.14292132	18.41619012	1.583977823	Up	0	0	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1
CCG007718.1	1227	175	738	6.12852776	21.0262496	1.778579098	Up	3.55E-13	9.58E-13	AAEL002532-PA OS=Aedes aegypti GN=AAEL002532 PE=4 SV=1
CCG007631.1	894	186	738	8.9400119	28.85817479	1.690631399	Up	6.43E-13	1.64E-12	AAEL007492-PA OS=Aedes aegypti GN=AAEL007492 PE=4 SV=1
CCG018215.1	1559	281	738	7.74502593	16.54856206	1.09536389	Up	0	0	AAEL001190-PA OS=Aedes aegypti GN=AAEL001190 PE=4 SV=1
CCG013917.1	2205	32	739	0.62359706	11.7161753	4.231743758	Up	1.77E-13	5.17E-13	AAEL002810-PA OS=Aedes aegypti GN=AAEL002810 PE=3 SV=1
CCG004103.1	3621	61	739	0.72387567	7.134539225	3.30100642	Up	1.75E-14	6.18E-14	AGAP007129-PA OS=Anopheles gambiae GN=AGAP007129 PE=4 SV=4
CCG025749.1	1146	98	739	3.67454973	22.54290274	2.617033914	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_13112 PE=4 SV=1
CCG022733.1	1073	109	739	4.36505226	24.07657645	2.463559433	Up	0	0	AAEL006986-PA (Fragment) OS=Aedes aegypti GN=AAEL006986 PE=4 SV=1
CCG005478.1	861	244	739	12.1772535	30.00483918	1.30100642	Up	2.78E-13	7.73E-13	AAEL010071-PA OS=Aedes aegypti GN=AAEL010071 PE=3 SV=1
CCG027701.1	1792	188	740	4.50798555	14.43589554	1.679105813	Up	1.80E-13	5.25E-13	AAEL002097-PA OS=Aedes aegypti GN=AAEL002097 PE=4 SV=1
CCG011774.1	1853	237	740	5.4958592	13.96067178	1.344951415	Up	2.32E-13	6.61E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011988 PE=4 SV=1
CCG017631.1	2346	295	740	5.40327012	11.02690742	1.02912352	Up	2.32E-13	6.62E-13	AAEL004986-PA (Fragment) OS=Aedes aegypti GN=AAEL004986 PE=4 SV=1

CCG025080.1	697	55	741	3.39072511	37.16511203	3.454283223	Up	0	0	AAEL001402-PA OS=Aedes aegypti GN=AAEL001402 PE=4 SV=1
CCG027357.1	2619	86	742	1.41099549	9.904177688	2.811323826	Up	0	0	AAEL004896-PA OS=Aedes aegypti GN=AAEL004896 PE=4 SV=1
CCG027736.1	473	122	742	11.0831028	54.83941092	2.306851243	Up	8.04E-14	2.58E-13	AAEL001623-PA OS=Aedes aegypti GN=AAEL001623 PE=2 SV=1
CCG003087.1	1965	259	742	5.6636953	13.20052996	1.220780293	Up	0	0	AAEL005149-PA (Fragment) OS=Aedes aegypti GN=AAEL005149 PE=4 SV=1
CCG022626.1	2152	1315	742	26.2570637	12.05345788	-1.123258504	Down	1.36E-67	6.68E-67	AAEL001104-PA OS=Aedes aegypti GN=AAEL001104 PE=4 SV=1
CCG027176.1	1012	1436	742	60.9728645	25.6314638	-1.250251453	Down	4.35E-87	2.24E-86	Putative short-chain dehydrogenase/reductase Mc1 OS=Aedes aegypti GN=SDR2 PE=2 SV=1
CCG000044.1	960	2708	742	121.21046	27.01983476	-2.165423443	Down	0	0	AAEL006953-PA OS=Aedes aegypti GN=AAEL006953 PE=4 SV=1
CCG020766.1	1981	101	743	2.19078405	13.11155964	2.581320122	Up	0	0	AAEL015455-PA OS=Aedes aegypti GN=AAEL015455 PE=4 SV=1
CCG008303.1	2283	129	743	2.42798763	11.37713519	2.228304349	Up	0	0	Zinc finger protein 109 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004616 PE=4 SV=1
CCG018887.2	2719	175	743	2.76561367	9.552776624	1.788320493	Up	3.55E-13	9.57E-13	AAEL011413-PA OS=Aedes aegypti GN=AAEL011413 PE=4 SV=1
CCG011473.1	1773	266	743	6.44667197	14.64974599	1.184249169	Up	4.09E-14	1.40E-13	AAEL002115-PA OS=Aedes aegypti GN=AAEL002115 PE=4 SV=1
CCG025953.1	1401	1559	743	47.8157147	18.53961431	-1.366873608	Down	3.27E-108	1.75E-107	AAEL013654-PA OS=Aedes aegypti GN=AAEL013654 PE=4 SV=1
CCG002862.1	852	2069	743	104.347865	30.48591507	-1.775186326	Down	1.77E-209	1.06E-208	AAEL008601-PA OS=Aedes aegypti GN=AAEL008601 PE=2 SV=1
CCG006133.1	1485	8	744	0.23148679	17.51444978	6.241472015	Up	2.19E-06	3.97E-06	AAEL006209-PA OS=Aedes aegypti GN=AAEL006209 PE=4 SV=1
CCG005152.1	3024	98	744	1.3925377	8.600845873	2.626762171	Up	0	0	histone-lysine N-methyltransferase ash1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001202 PE=4 SV=1

CCG017466.1	1225	114	744	3.99881612	21.23180238	2.408582001	Up	8.48E-14	2.70E-13	AAEL009476-PA OS=Aedes aegypti GN=AAEL009476 PE=4 SV=1
CCG010072.1	2786	114	744	1.75827342	9.3355915	2.408582001	Up	8.48E-14	2.70E-13	AAEL006491-PA OS=Aedes aegypti GN=AAEL006491 PE=4 SV=1
CCG028042.1	2746	198	744	3.09832755	9.471579723	1.612115395	Up	0	0	XP4-binding protein Z OS=Culex quinquefasciatus GN=CpipJ_CPIJ005885 PE=4 SV=1
CCG021665.1	3159	280	744	3.80865011	8.233288357	1.112188998	Up	0	0	AAEL013108-PA OS=Aedes aegypti GN=AAEL013108 PE=4 SV=1
CCG018772.1	1998	1788	744	38.4533961	13.01749646	-1.562659006	Down	3.82E-151	2.17E-150	AAEL010047-PA OS=Aedes aegypti GN=AAEL010047 PE=3 SV=1
CCG025139.1	521	1976	744	162.971585	49.92122441	-1.706895217	Down	2.03E-189	1.20E-188	Salivary cysteine-rich peptide OS=Culex quinquefasciatus GN=CpipJ_CPIJ006214 PE=4 SV=1
CCG014400.1	2046	85	745	1.78515515	12.7291868	2.834018883	Up	0	0	AAEL005452-PA OS=Aedes aegypti GN=AAEL005452 PE=4 SV=1
CCG022180.1	1896	109	746	2.47030647	13.75468063	2.477160699	Up	0	0	AAEL007208-PA OS=Aedes aegypti GN=AAEL007208 PE=4 SV=1
CCG017889.2	2412	110	746	1.95964793	10.81213701	2.463985311	Up	0	0	AAEL013590-PA OS=Aedes aegypti GN=AAEL013590 PE=4 SV=1
CCG001238.1	720	300	746	17.9040561	36.22065899	1.016526334	Up	1.10E-13	3.42E-13	AAEL000939-PA OS=Aedes aegypti GN=AAEL000939 PE=4 SV=1
CCG017056.1	255	2446	746	412.172435	102.270096	-2.010863664	Down	5.40E-293	3.39E-292	AAEL008802-PA OS=Aedes aegypti GN=AAEL008802 PE=4 SV=1
CCG002482.1	1254	7	747	0.23986295	20.82442803	6.439922715	Up	4.31E-06	7.70E-06	AAEL006884-PA OS=Aedes aegypti GN=AAEL006884 PE=4 SV=1
CCG005515.1	1757	62	747	1.51629115	14.86273919	3.293081326	Up	0	0	AAEL006177-PA OS=Aedes aegypti GN=AAEL006177 PE=4 SV=1
CCG027119.1	1848	93	747	2.16243794	14.13086188	2.708118825	Up	4.04E-14	1.39E-13	AAEL010986-PA (Fragment) OS=Aedes aegypti GN=AAEL010986 PE=3 SV=1
CCG025587.1	1938	126	747	2.79369792	13.4746299	2.269997713	Up	1.34E-13	4.06E-13	AAEL002136-PA OS=Aedes aegypti GN=AAEL002136 PE=4 SV=1
CCG021335.1	1128	144	747	5.48549804	23.15056095	2.077352635	Up	7.53E-14	2.44E-13	AAEL013389-PA OS=Aedes aegypti GN=AAEL013389 PE=4 SV=1

CCG021819.1	2514	247	747	4.22176788	10.38736386	1.298910405	Up	0	0	AAEL010569-PA OS=Aedes aegypti GN=AAEL010569 PE=4 SV=1
CCG024445.1	2788	2463	747	37.9607089	9.366511029	-2.018923276	Down	1.40E-296	8.81E-296	AAEL006240-PA OS=Aedes aegypti GN=AAEL006240 PE=4 SV=1
CCG008821.1	300	8	748	1.14585959	87.16263675	6.249207664	Up	2.19E-06	3.97E-06	Defensin OS=Culex quinquefasciatus PE=4 SV=1
CCG023463.1	1602	39	748	1.04607968	16.32259115	3.963805445	Up	7.33E-14	2.38E-13	AAEL003878-PA OS=Aedes aegypti GN=AAEL003878 PE=4 SV=1
CCG008013.1	1023	288	748	12.0970514	25.56089054	1.079282662	Up	0	0	Methylthioribose-1-phosphate isomerase OS=Glossina morsitans morsitans PE=2 SV=1
CCG023669.1	1204	52	749	1.85583571	21.74730009	3.550695394	Up	0	0	AAEL002045-PA OS=Aedes aegypti GN=AAEL002045 PE=4 SV=1
CCG005845.1	1875	173	749	3.96467418	13.96466629	1.816506885	Up	3.60E-13	9.68E-13	Mitochondrial GTPase 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017444 PE=3 SV=1
CCG028506.1	1152	222	749	8.28062594	22.72894905	1.456719246	Up	0	0	AAEL000463-PA OS=Aedes aegypti GN=AAEL000463 PE=4 SV=1
CCG015331.1	1326	1591	749	51.5572004	19.74641727	-1.384583008	Down	1.45E-112	7.83E-112	AAEL004137-PA OS=Aedes aegypti GN=AAEL004137 PE=3 SV=1
CCG002829.1	1965	17	750	0.37174834	13.34285373	5.165597148	Up	3.39E-09	6.88E-09	AAEL002281-PA OS=Aedes aegypti GN=AAEL002281 PE=4 SV=1
CCG024403.1	1731	74	750	1.83694995	15.14656706	3.043606624	Up	0	0	AAEL002751-PA OS=Aedes aegypti GN=AAEL002751 PE=4 SV=1
CCG006027.1	1519	92	750	2.60251191	17.26050532	2.729498033	Up	1.44E-14	5.12E-14	AAEL012931-PA OS=Aedes aegypti GN=AAEL012931 PE=4 SV=1
CCG011073.1	984	170	750	7.42363301	26.64502803	1.843669053	Up	3.53E-13	9.52E-13	Short-chain dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ009166 PE=3 SV=1
CCG021023.1	4213	256	750	2.61102589	6.223286869	1.253059989	Up	0	0	AAEL006226-PA OS=Aedes aegypti GN=AAEL006226 PE=4 SV=1
CCG012256.1	678	69	751	4.37302609	38.72222103	3.146457845	Up	0	0	GM16159 OS=Drosophila sechellia GN=GM16159 PE=4 SV=1

CCG011183.1	1140	293	751	11.0439757	23.02953145	1.060225447	Up	4.19E-13	1.11E-12	StAR-related lipid transfer domain containing protein OS=Aedes aegypti PE=2 SV=1
CCG014159.1	2871	296	751	4.43017814	9.144432552	1.045528936	Up	2.26E-13	6.47E-13	AAEL001581-PA OS=Aedes aegypti GN=AAEL001581 PE=4 SV=1
CCG024801.1	1037	296	751	12.265228	25.31693911	1.045528936	Up	2.26E-13	6.47E-13	AAEL000421-PA OS=Aedes aegypti GN=AAEL000421 PE=4 SV=1
CCG016003.1	2081	305	751	6.29782271	12.61588941	1.002316869	Up	5.30E-13	1.37E-12	AAEL002329-PB OS=Aedes aegypti GN=AAEL002329 PE=4 SV=1
CCG009653.1	879	107	752	5.23067304	29.90742222	2.515435069	Up	0	0	AAEL013942-PA OS=Aedes aegypti GN=AAEL013942 PE=4 SV=1
CCG020515.1	3075	127	752	1.77468497	8.549146059	2.268217369	Up	1.57E-13	4.66E-13	AAEL010600-PA OS=Aedes aegypti GN=AAEL010600 PE=4 SV=1
CCG019494.1	1056	101	753	4.10979469	24.92763486	2.600607776	Up	0	0	AAEL004285-PA OS=Aedes aegypti GN=AAEL004285 PE=4 SV=1
CCG014271.1	1389	101	753	3.12450914	18.95146322	2.600607776	Up	0	0	AAEL012211-PA OS=Aedes aegypti GN=AAEL012211 PE=4 SV=1
CCG017114.1	1239	107	753	3.71086489	21.24582922	2.517352272	Up	0	0	AAEL000502-PA (Fragment) OS=Aedes aegypti GN=AAEL000502 PE=4 SV=1
CCG006657.1	1338	161	753	5.17049871	19.67382841	1.92790238	Up	0	0	AAEL002159-PA OS=Aedes aegypti GN=AAEL002159 PE=4 SV=1
CCG002570.1	836	286	753	14.7001724	31.48753877	1.098947922	Up	0	0	AAEL004570-PA OS=Aedes aegypti GN=AAEL004570 PE=4 SV=1
CCG016373.1	3896	1530	753	16.8746648	6.756566327	-1.320496679	Down	8.48E-101	4.50E-100	AAEL005763-PA (Fragment) OS=Aedes aegypti GN=AAEL005763 PE=4 SV=1
CCG023039.1	1863	154	754	3.55198021	14.14843837	1.993947376	Up	1.82E-13	5.29E-13	JmjC domain-containing histone demethylation protein 3C OS=Culex quinquefasciatus GN=CpipJ_CPIJ011492 PE=4 SV=1
CCG002218.1	1791	225	754	5.39820787	14.71721981	1.446952726	Up	0	0	AAEL002828-PB (Fragment) OS=Aedes aegypti GN=AAEL002828 PE=4 SV=1

CCG017290.1	1697	1485	754	37.6016829	15.53243411	-1.275513299	Down	8.07E-93	4.22E-92	AAEL014762-PA OS=Aedes aegypti GN=AAEL014762 PE=4 SV=1
CCG021032.1	2148	2099	754	41.9895125	12.27120144	-1.774752534	Down	2.06E-212	1.24E-211	Uncharacterized protein OS=Anopheles darlingi GN=AND_18243 PE=4 SV=1
CCG007045.1	1787	2598	754	62.4708285	14.75016267	-2.082451798	Down	0	0	Galactose-1-phosphate uridylyltransferase OS=Aedes aegypti GN=AAEL001607 PE=3 SV=1
CCG022950.1	1735	81	755	2.00607983	15.21239133	2.922796035	Up	4.46E-14	1.51E-13	AAEL004945-PA (Fragment) OS=Aedes aegypti GN=AAEL004945 PE=4 SV=1
CCG011503.1	2703	122	755	1.93944048	9.764520519	2.3319087	Up	8.04E-14	2.58E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015878 PE=4 SV=1
CCG006917.1	1105	152	755	5.91076892	23.88551942	2.014718525	Up	8.39E-14	2.69E-13	AAEL009356-PA OS=Aedes aegypti GN=AAEL009356 PE=4 SV=1
CCG018662.1	1328	154	755	4.98293609	19.87462271	1.995859497	Up	1.82E-13	5.28E-13	AAEL014061-PA (Fragment) OS=Aedes aegypti GN=AAEL014061 PE=4 SV=1
CCG025205.1	1579	155	755	4.21805501	16.7153255	1.986521633	Up	3.42E-13	9.26E-13	AAEL007811-PA OS=Aedes aegypti GN=AAEL007811 PE=2 SV=1
CCG019859.1	18463	212	755	0.49339673	1.429534689	1.534725583	Up	0	0	AAEL007993-PA OS=Aedes aegypti GN=CLIPB27 PE=3 SV=1
CCG024607.1	315	1617	755	220.577971	83.7888856	-1.396457925	Down	7.42E-116	4.02E-115	W-rich putative salivary secreted peptide OS=Aedes albopictus PE=4 SV=1
CCG020869.1	3429	34	756	0.42606328	7.707336611	4.177092787	Up	8.82E-14	2.80E-13	AAEL000201-PA (Fragment) OS=Aedes aegypti GN=AAEL000201 PE=4 SV=1
CCG019378.3	3936	36	756	0.39301587	6.714547063	4.094630627	Up	6.13E-14	2.03E-13	AAEL006658-PA OS=Aedes aegypti GN=AAEL006658 PE=4 SV=1
CCG018690.1	651	189	756	12.4750842	40.59670851	1.702313204	Up	2.34E-13	6.64E-13	AAEL004632-PA OS=Aedes aegypti GN=AAEL004632 PE=4 SV=1

CCG009005.1	1836	129	757	3.01911534	14.41362501	2.255235438	Up	0	0	Lamin OS=Culex quinquefasciatus GN=CpipJ_CPIJ014365 PE=4 SV=1
CCG022387.1	1296	1756	758	58.2213379	20.44627608	-1.509709888	Down	3.74E-141	2.10E-140	AAEL000516-PA OS=Aedes aegypti GN=AAEL000516 PE=4 SV=1
CCG004239.3	2730	2035	758	32.0305531	9.706364027	-1.722445837	Down	1.57E-197	9.36E-197	AAEL007024-PA OS=Aedes aegypti GN=CYP6AG3 PE=3 SV=1
CCG013319.1	804	2156	758	115.227298	32.95817636	-1.805774221	Down	1.90E-223	1.15E-222	AAEL007659-PA OS=Aedes aegypti GN=AAEL007659 PE=4 SV=1
CCG012464.1	1534	37	759	1.03642776	17.29682664	4.060815914	Up	5.17E-14	1.74E-13	AAEL001785-PA OS=Aedes aegypti GN=AAEL001785 PE=4 SV=1
CCG011158.1	2982	44	759	0.63402694	8.897831009	3.810837661	Up	3.91E-14	1.35E-13	Riepper OS=Culex quinquefasciatus GN=CpipJ_CPIJ004355 PE=4 SV=1
CCG024034.1	1770	64	759	1.55370792	14.99058309	3.270269279	Up	0	0	AAEL004654-PA OS=Aedes aegypti GN=AAEL004654 PE=4 SV=1
CCG004803.1	1078	91	759	3.62731526	24.61348058	2.762474639	Up	0	0	Hybrid hormone receptor- associated protein 6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004789 PE=4 SV=1
CCG008517.1	5348	124	759	0.99630649	4.961356034	2.316072969	Up	1.61E-13	4.76E-13	AAEL010698-PA (Fragment) OS=Aedes aegypti GN=AAEL010698 PE=4 SV=1
CCG009856.1	1179	2	760	0.07289183	22.53459741	8.272168812	Up	9.36E-05	0.00015399	AAEL012453-PA OS=Aedes aegypti GN=AAEL012453 PE=4 SV=1
CCG017182.1	2154	93	760	1.85523924	12.33439663	2.733010001	Up	4.04E-14	1.38E-13	AAEL013590-PA OS=Aedes aegypti GN=AAEL013590 PE=4 SV=1
CCG023973.1	1011	171	760	7.26787796	26.27921894	1.854316297	Up	3.42E-13	9.27E-13	AAEL007015-PA OS=Aedes aegypti GN=AAEL007015 PE=4 SV=1
CCG014863.1	2559	200	760	3.35832236	10.382294	1.628312622	Up	0	0	AAEL008539-PA OS=Aedes aegypti GN=AAEL008539 PE=4 SV=1
CCG025490.1	2582	241	760	4.01073046	10.28981036	1.359279476	Up	2.37E-13	6.71E-13	AAEL000853-PA OS=Aedes aegypti GN=AAEL000853 PE=4 SV=1
CCG026698.1	613	257	760	18.0150437	43.34141982	1.266544263	Up	4.14E-13	1.10E-12	AAEL000983-PA OS=Aedes aegypti GN=AAEL000983 PE=4 SV=1

CCG010785.2	2333	280	760	5.15710488	11.38803701	1.142885795	Up	0	0	AAEL007151-PA OS=Aedes aegypti GN=AAEL007151 PE=4 SV=1
CCG003816.1	3243	3333	760	44.1622342	8.192503961	-2.43043679	Down	0	0	AAEL008635-PA OS=Aedes aegypti GN=AAEL008635 PE=3 SV=1
CCG002424.1	273	89	761	14.0084483	97.44779716	2.798332416	Up	0	0	AAEL006830-PA OS=Aedes aegypti GN=AAEL006830 PE=4 SV=1
CCG017174.1	951	95	761	4.29245509	27.97397332	2.704210239	Up	0	0	AAEL011301-PA OS=Aedes aegypti GN=AAEL011301 PE=4 SV=1
CCG027540.1	1524	227	761	6.40034761	17.45619988	1.44751736	Up	0	0	AAEL015415-PA OS=Aedes aegypti GN=AAEL015415 PE=4 SV=1
CCG000033.1	3494	304	761	3.73863747	7.613980717	1.026138334	Up	0	0	Peptidase OS=Culex quinquefasciatus GN=CpipJ_CPIJ014636 PE=4 SV=1
CCG027127.1	1489	306	761	8.83058347	17.86652023	1.016678005	Up	0	0	Ultraspiracle OS=Aedes albopictus GN=USP PE=2 SV=1
CCG025903.1	1866	2165	761	49.8550243	14.25683206	-1.806085462	Down	2.00E-224	1.21E-223	AAEL004567-PA OS=Aedes aegypti GN=AAEL004567 PE=4 SV=1
CCG019181.1	1622	8	762	0.21193457	16.42306221	6.275960391	Up	2.19E-06	3.97E-06	Cytochrome P450 6CB1 (Fragment) OS=Aedes aegypti GN=CYP6CB1 PE=2 SV=1
CCG009203.1	1134	85	762	3.22083549	23.49048228	2.866569455	Up	0	0	AAEL003124-PA OS=Aedes aegypti GN=AAEL003124 PE=4 SV=1
CCG002269.1	2162	121	762	2.40487414	12.32109477	2.357097154	Up	0	0	AAEL003273-PA OS=Aedes aegypti GN=AAEL003273 PE=4 SV=1
CCG017522.1	2508	196	762	3.35808133	10.62129462	1.661250547	Up	7.93E-14	2.55E-13	Glutamyl-tRNA(Gln) amidotransferase subunit A, mitochondrial OS=Drosophila mojavensis GN=GI22341 PE=3 SV=1
CCG024784.1	1576	280	762	7.63421681	16.90241555	1.146677374	Up	0	0	AAEL011912-PA OS=Aedes aegypti GN=AAEL011912 PE=2 SV=1
CCG018676.1	1816	77	764	1.82195461	14.70711644	3.012955491	Up	0	0	AAEL009786-PA OS=Aedes aegypti GN=AAEL009786 PE=4 SV=1

CCG011675.1	1296	95	764	3.14978765	20.60811995	2.709886424	Up	0	0	AAEL004371-PA (Fragment) OS=Aedes aegypti GN=AAEL004371 PE=4 SV=1
CCG012326.1	1516	244	764	6.91597312	17.61749568	1.349004694	Up	2.78E-13	7.73E-13	UDP-galactose transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ007457 PE=4 SV=1
CCG010047.1	1131	1736	764	65.9553132	23.6146096	-1.481809201	Down	9.02E-136	5.04E-135	-
CCG021397.1	6170	24	765	0.16714321	4.334373052	4.696666641	Up	1.93E-11	4.24E-11	Polyprotein OS=Antheraea mylitta PE=4 SV=1
CCG015241.1	912	28	765	1.31924624	29.32355453	4.474274219	Up	1.17E-12	2.85E-12	AAEL010327-PA OS=Aedes aegypti GN=AAEL010327 PE=3 SV=1
CCG008031.1	3603	248	765	2.95767255	7.42244844	1.327432831	Up	5.86E-14	1.95E-13	Aluminate kinase SKIZ OS=Culex quinquefasciatus GN=CpipJ_CPIJ003293 PE=4 SV=1
CCG002999.1	1216	287	765	10.1417055	21.9926659	1.116722215	Up	2.87E-13	7.99E-13	AAEL014050-PA OS=Aedes aegypti GN=AAEL014050 PE=4 SV=1
CCG021783.1	1671	18	766	0.46286967	16.02515859	5.113588784	Up	1.62E-09	3.33E-09	AAEL009057-PA OS=Aedes aegypti GN=AAEL009057 PE=3 SV=1
CCG026554.1	1495	49	766	1.40837257	17.91173245	3.668803942	Up	0	0	AAEL011838-PA OS=Aedes aegypti GN=AAEL011838 PE=4 SV=1
CCG021616.1	1859	72	766	1.66423932	14.40454008	3.113588784	Up	0	0	Cell division protein kinase 6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000891 PE=4 SV=1
CCG022837.1	1341	295	766	9.45270075	19.96870992	1.078942641	Up	2.32E-13	6.62E-13	Spastin OS=Culex quinquefasciatus GN=CpipJ_CPIJ013667 PE=3 SV=1
CCG003046.1	2214	305	766	5.91949822	12.09486902	1.030848353	Up	5.30E-13	1.37E-12	AAEL010404-PA OS=Aedes aegypti GN=AAEL010404 PE=4 SV=1
CCG021962.1	18936	1477	766	3.35162115	1.414133925	-1.244940325	Down	6.22E-89	3.23E-88	AAEL010769-PA OS=Aedes aegypti GN=SRPN6 PE=3 SV=1
CCG008081.1	4440	1631	766	15.784603	6.031090092	-1.388027281	Down	8.84E-116	4.80E-115	AAEL000564-PA OS=Aedes aegypti GN=AAEL000564 PE=4 SV=1
CCG019819.1	3219	176	767	2.34938592	8.329604935	1.825964353	Up	0	0	AAEL009289-PA OS=Aedes aegypti GN=AAEL009289 PE=4 SV=1
CCG028570.1	2209	192	767	3.73480717	12.13807075	1.700433471	Up	0	0	AAEL006459-PA OS=Aedes aegypti GN=AAEL006459 PE=4 SV=1

CCG013655.1	6722	270	767	1.72594888	3.988842351	1.208580374	Up	2.59E-13	7.26E-13	AAEL010461-PA OS=Aedes aegypti GN=AAEL010461 PE=4 SV=1
CCG016411.1	1647	1250	767	32.612124	16.27990181	-1.002316408	Down	1.16E-53	5.50E-53	AAEL010348-PA OS=Aedes aegypti GN=AAEL010348 PE=3 SV=1
CCG013132.1	765	204	768	11.4585959	35.09536805	1.614850363	Up	6.59E-14	2.17E-13	AAEL009809-PA OS=Aedes aegypti GN=AAEL009809 PE=4 SV=1
CCG003004.1	1614	1784	768	47.495667	16.63442166	-1.513624195	Down	6.38E-144	3.60E-143	AAEL008672-PA OS=Aedes aegypti GN=AAEL008672 PE=3 SV=1
CCG004569.1	1612	190	769	5.06467095	16.67674618	1.719297383	Up	2.69E-13	7.53E-13	Cen division protein kinase 9 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008862 PE=4 SV=1
CCG011741.1	2040	203	769	4.27590987	13.17789943	1.623817075	Up	9.28E-14	2.93E-13	AAEL007387-PA OS=Aedes aegypti GN=AAEL007387 PE=4 SV=1
CCG015209.1	816	250	769	13.1647471	32.94474858	1.323368707	Up	2.10E-13	6.02E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003086 PE=4 SV=1
CCG020066.3	3699	89	770	1.03387574	7.277067617	2.815294408	Up	0	0	AAEL015044-PA OS=Aedes aegypti GN=AAEL015044 PE=4 SV=1
CCG025120.1	815	108	770	5.69414888	33.02806517	2.536140337	Up	0	0	AAEL005882-PA (Fragment) OS=Aedes aegypti GN=AAEL005882 PE=4 SV=1
CCG023747.3	1914	189	770	4.24309292	14.06367456	1.728785415	Up	2.34E-13	6.64E-13	AAEL009262-PA OS=Aedes aegypti GN=AAEL009262 PE=4 SV=1
CCG011418.1	3049	227	770	3.19912422	8.828426735	1.464479352	Up	0	0	AAEL010512-PA OS=Aedes aegypti GN=AAEL010512 PE=4 SV=1
CCG028236.1	1053	2846	770	116.136624	25.5630324	-2.183692107	Down	0	0	AGAP000969-PB OS=Anopheles gambiae GN=AgaP_AGAP000969 PE=4 SV=1
CCG007224.1	3045	54	771	0.76202485	8.851504562	3.538012752	Up	0	0	AAEL005047-PA OS=Aedes aegypti GN=AAEL005047 PE=4 SV=1
CCG019810.1	3169	75	772	1.01695491	8.516184812	3.065951551	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001516 PE=4 SV=1
CCG000213.1	1212	79	772	2.80083254	22.26715319	2.990989493	Up	0	0	AAEL003868-PA OS=Aedes aegypti GN=AAEL003868 PE=4 SV=1

CCG026640.1	2564	92	772	1.54181575	10.525659	2.771208285	Up	1.44E-14	5.11E-14	AAEL003170-PA OS=Aedes aegypti GN=AAEL003170 PE=4 SV=1
CCG016128.1	2937	94	772	1.3752656	9.188896721	2.740181389	Up	0	0	Transporter OS=Aedes aegypti GN=AAEL006800 PE=3 SV=1
CCG025407.1	2011	253	772	5.40593877	13.42008437	1.311776666	Up	3.13E-13	8.62E-13	AAEL014582-PA OS=Aedes aegypti GN=AAEL014582 PE=4 SV=1
CCG015826.1	1470	253	772	7.39547133	18.35904059	1.311776666	Up	3.13E-13	8.62E-13	AAEL002214-PA OS=Aedes aegypti GN=AAEL002214 PE=4 SV=1
CCG001117.1	1326	261	772	8.45784369	20.35278256	1.266864245	Up	8.02E-14	2.58E-13	AAEL000875-PA OS=Aedes aegypti GN=AAEL000875 PE=4 SV=1
CCG012733.1	13512	5318	772	16.9118597	1.997320135	-3.081897821	Down	0	0	AAEL008855-PA OS=Aedes aegypti GN=AAEL008855 PE=4 SV=1
CCG004494.1	1047	0	773	0.001	25.8096924	14.65562533	Up	4.44E-16	1.64E-15	AAEL007783-PA OS=Aedes aegypti GN=AAEL007783 PE=4 SV=1
CCG019030.1	1566	26	773	0.71341833	17.25590546	4.59619809	Up	4.60E-12	1.05E-11	AAEL010598-PA OS=Aedes aegypti GN=AAEL010598 PE=4 SV=1
CCG025891.1	3927	109	773	1.1926919	6.881270167	2.528453483	Up	0	0	AAEL010633-PA OS=Aedes aegypti GN=AAEL010633 PE=4 SV=1
CCG002085.1	2499	155	773	2.66518962	10.81342455	2.020513402	Up	3.42E-13	9.27E-13	AAEL007512-PA OS=Aedes aegypti GN=AAEL007512 PE=4 SV=1
CCG024057.1	2300	245	773	4.57721086	11.74902085	1.359999869	Up	3.40E-13	9.24E-13	AAEL013631-PA OS=Aedes aegypti GN=AAEL013631 PE=4 SV=1
CCG019308.1	1023	0	774	0.001	26.4493707	14.69094578	Up	4.44E-16	1.63E-15	AAEL013715-PA OS=Aedes aegypti GN=AAEL013715 PE=3 SV=1
CCG021609.1	1155	136	774	5.05963975	23.42658547	2.211040119	Up	1.70E-13	4.98E-13	AAEL008391-PA OS=Aedes aegypti GN=AAEL008391 PE=4 SV=1
CCG011103.1	357	227	774	27.3224923	75.79189418	1.471954473	Up	0	0	AAEL003764-PA OS=Aedes aegypti GN=AAEL003764 PE=4 SV=1
CCG004807.1	1314	229	774	7.48863716	20.59186166	1.459299172	Up	2.46E-13	6.92E-13	AAEL007885-PA OS=Aedes aegypti GN=AAEL007885 PE=4 SV=1
CCG016125.1	1170	2	775	0.07345254	23.1561235	8.300365704	Up	9.36E-05	0.00015361	AAEL006726-PA OS=Aedes aegypti GN=AAEL006726 PE=4 SV=1
CCG023393.1	3780	155	775	1.76198647	7.16737156	2.024241299	Up	3.42E-13	9.28E-13	AAEL009294-PA OS=Aedes aegypti GN=AAEL009294 PE=4 SV=1

CCG027036.3	1770	67	776	1.62653798	15.32634055	3.236136856	Up	0	0	AAEL006338-PA OS=Aedes aegypti GN=AAEL006338 PE=2 SV=1
CCG002276.1	1170	8	777	0.29381015	23.21588124	6.304083992	Up	2.19E-06	3.96E-06	AAEL006726-PA OS=Aedes aegypti GN=AAEL006726 PE=4 SV=1
CCG007183.1	3129	50	777	0.68663686	8.680914366	3.660227802	Up	0	0	AAEL007123-PA (Fragment) OS=Aedes aegypti GN=AAEL007123 PE=3 SV=2
CCG019687.1	1254	112	777	3.83780724	21.66075044	2.49672907	Up	0	0	AAEL005582-PA OS=Aedes aegypti GN=AAEL005582 PE=4 SV=1
CCG013804.1	1333	167	778	5.38330509	20.40325531	1.922235256	Up	1.62E-13	4.78E-13	AAEL009971-PA OS=Aedes aegypti GN=AAEL009971 PE=4 SV=1
CCG000477.1	1294	77	779	2.55693166	21.0452068	3.041006181	Up	0	0	AAEL000130-PA OS=Aedes aegypti GN=AAEL000130 PE=4 SV=1
CCG009641.1	1512	98	779	2.78507539	18.01091111	2.693082878	Up	0	0	AAEL014868-PA OS=Aedes aegypti GN=AAEL014868 PE=4 SV=1
CCG001043.1	1548	166	779	4.6078656	17.59205272	1.932753291	Up	1.83E-13	5.32E-13	AAEL000262-PA OS=Aedes aegypti GN=AAEL000262 PE=4 SV=1
CCG011354.1	551	1	780	0.077985	49.48721576	9.309643518	Up	0.000587	0.00091751	-
CCG017483.1	948	16	780	0.72522759	28.76313912	5.309643518	Up	7.09E-09	1.42E-08	Putative salivary protein 34k-1 OS=Aedes albopictus PE=2 SV=1
CCG014024.1	465	106	780	9.79525133	58.63969007	2.581723063	Up	0	0	AAEL003469-PA OS=Aedes aegypti GN=AAEL003469 PE=4 SV=1
CCG001404.1	1141	108	780	4.0672492	23.89785792	2.554756015	Up	0	0	AAEL012040-PA OS=Aedes aegypti GN=AAEL012040 PE=4 SV=1
CCG012661.1	1875	116	780	2.65839425	14.54264314	2.451662522	Up	1.16E-13	3.59E-13	AAEL011220-PA OS=Aedes aegypti GN=AAEL011220 PE=4 SV=1
CCG026268.1	2046	131	780	2.75123912	13.32720229	2.276220516	Up	1.26E-13	3.87E-13	AAEL002973-PA OS=Aedes aegypti GN=AAEL002973 PE=4 SV=1
CCG009962.1	4482	137	780	1.31344347	6.083769719	2.211611435	Up	2.90E-13	8.04E-13	AAEL014774-PA OS=Aedes aegypti GN=AAEL014774 PE=4 SV=1
CCG020799.1	1248	92	781	3.16764069	21.87693442	2.787929986	Up	1.44E-14	5.12E-14	AAEL009474-PA OS=Aedes aegypti GN=PGRPS1 PE=4 SV=2
CCG021986.1	1229	150	781	5.24447534	22.21514578	2.082673251	Up	0	0	-
CCG022071.1	4191	180	781	1.84551473	6.514534517	1.819638846	Up	3.32E-13	9.05E-13	AAEL004895-PA OS=Aedes aegypti GN=AAEL004895 PE=4 SV=1

CCG006754.1	2457	262	781	4.58203926	11.11209367	1.27806894	Up	0	0	AAEL004551-PA OS=Aedes aegypti GN=AAEL004551 PE=4 SV=1
CCG008934.1	6261	1362	781	9.34751294	4.360711413	-1.100019046	Down	1.15E-67	5.67E-67	AAEL011241-PA OS=Aedes aegypti GN=AAEL011241 PE=4 SV=1
CCG006377.1	2781	96	782	1.48331338	9.830051218	2.7283755	Up	0	0	AAEL002496-PA OS=Aedes aegypti GN=AAEL002496 PE=4 SV=1
CCG008661.1	737	102	782	5.94696463	37.09277128	2.640912659	Up	0	0	GH15970 OS=Drosophila grimshawi GN=GH15970 PE=4 SV=1
CCG011667.1	2379	221	782	3.99172398	11.49111914	1.525435442	Up	0	0	AAEL005441-PA (Fragment) OS=Aedes aegypti GN=AAEL005441 PE=4 SV=1
CCG022502.1	1227	1554	782	54.4213265	22.27984714	-1.288432787	Down	1.46E-98	7.71E-98	AAEL000328-PA OS=Aedes aegypti GN=AAEL000328 PE=4 SV=1
CCG007341.1	1740	21	783	0.51860025	15.73122455	4.922864278	Up	1.76E-10	3.73E-10	AAEL009057-PA OS=Aedes aegypti GN=AAEL009057 PE=3 SV=1
CCG009362.1	2880	168	783	2.50656785	9.504281497	1.922864278	Up	1.45E-13	4.33E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007254 PE=4 SV=1
CCG002274.1	4755	244	783	2.2049664	5.756536427	1.384444364	Up	2.78E-13	7.73E-13	AAEL002818-PA OS=Aedes aegypti GN=AAEL002818 PE=4 SV=1
CCG027131.1	1789	260	783	6.24490274	15.30035255	1.292813888	Up	3.22E-13	8.81E-13	AGAP002097-PA OS=Anopheles gambiae GN=AgaP_AGAP002097 PE=4 SV=5
CCG009662.1	1715	2693	783	67.4737582	15.96054269	-2.079816813	Down	0	0	AAEL010951-PA OS=Aedes aegypti GN=AAEL010951 PE=3 SV=1
CCG024984.1	1476	55	784	1.60117575	18.56862398	3.535663334	Up	0	0	AAEL012260-PA OS=Aedes aegypti GN=AAEL012260 PE=4 SV=1
CCG002651.1	1439	108	784	3.22496966	19.04606601	2.562135546	Up	0	0	AAEL011317-PA OS=Aedes aegypti GN=AAEL011317 PE=3 SV=1
CCG003748.1	708	108	784	6.55470528	38.71086015	2.562135546	Up	0	0	AAEL011627-PA OS=Aedes aegypti GN=AAEL011627 PE=4 SV=1
CCG018310.1	1227	283	784	9.91070489	22.33682884	1.172364805	Up	0	0	Discs overgrown protein kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ003503 PE=3 SV=1

CCG013589.1	1479	38	785	1.10402293	18.55459585	4.070934534	Up	4.26E-14	1.45E-13	AAEL012520-PA OS=Aedes aegypti GN=AAEL012520 PE=4 SV=1
CCG013154.1	1015	72	785	3.0480994	27.03669681	3.148937046	Up	0	0	AAEL004482-PA OS=Aedes aegypti GN=AAEL004482 PE=4 SV=1
CCG017099.1	1788	214	785	5.14291007	15.34801301	1.577395061	Up	0	0	AAEL006306-PA OS=Aedes aegypti GN=AAEL006306 PE=4 SV=1
CCG018709.1	2760	233	785	3.62751745	9.942843212	1.454675903	Up	0	0	Paq OS=Culex quinquefasciatus GN=CpipJ_CPIJ018849 PE=4 SV=1
CCG024146.1	4818	270	785	2.40801751	5.695775688	1.242046451	Up	2.59E-13	7.26E-13	AAEL001142-PA OS=Aedes aegypti GN=AAEL001142 PE=4 SV=1
CCG023212.1	1527	304	785	8.55455096	17.97134726	1.070934534	Up	0	0	AAEL000415-PA OS=Aedes aegypti GN=AAEL000415 PE=4 SV=1
CCG020321.1	1343	28	786	0.89586937	20.45957226	4.513343784	Up	1.17E-12	2.85E-12	AAEL011261-PA OS=Aedes aegypti GN=AAEL011261 PE=4 SV=1
CCG021944.1	1770	140	786	3.39873607	15.52384494	2.191415689	Up	0	0	AAEL002427-PA OS=Aedes aegypti GN=AAEL002427 PE=4 SV=1
CCG021772.1	2040	215	786	4.52867301	13.4692184	1.572505856	Up	0	0	AAEL000003-PA OS=Aedes aegypti GN=AAEL000003 PE=4 SV=1
CCG000071.1	1710	130	787	3.26670497	16.08898469	2.300165216	Up	0	0	Zinc finger protein 569 OS=Culex quinquefasciatus GN=CpipJ_CPIJ020006 PE=4 SV=1
CCG004783.1	1139	238	787	8.97875052	24.15466534	1.427715266	Up	6.86E-13	1.73E-12	G73-specific cyclin-C OS=Culex quinquefasciatus GN=CpipJ_CPIJ006239 PE=3 SV=1
CCG016288.1	1027	273	787	11.4223345	26.78886448	1.229775888	Up	1.53E-14	5.42E-14	AAEL006570-PA OS=Aedes aegypti GN=AAEL006570 PE=4 SV=1
CCG005061.1	13465	302	787	0.96374748	2.043235338	1.08412829	Up	4.59E-13	1.20E-12	Salivary purine nucleosidase OS=Aedes albopictus PE=2 SV=1
CCG023004.1	975	0	788	0.001	28.25345856	14.78613986	Up	4.44E-16	1.63E-15	AAEL008137-PA OS=Aedes aegypti GN=AAEL008137 PE=4 SV=1
CCG000746.1	1563	77	788	2.11687112	17.6245183	3.057578483	Up	0	0	AAEL010437-PA (Fragment) OS=Aedes aegypti GN=AAEL010437 PE=4 SV=1
CCG022475.1	5124	218	788	1.8281425	5.376097209	1.556180698	Up	0	0	Pol-like protein OS=Biomphalaria glabrata PE=4 SV=1

CCG011411.1	777	229	788	12.6641818	35.45318159	1.485161235	Up	2.46E-13	6.92E-13	AAEL009634-PA OS=Aedes aegypti GN=AAEL009634 PE=3 SV=1
CCG026021.1	1821	1482	788	34.9704265	15.12746958	-1.208964709	Down	2.99E-85	1.54E-84	AAEL003586-PA OS=Aedes aegypti GN=AAEL003586 PE=4 SV=1
CCG023940.1	4203	1681	788	17.1858491	6.554157054	-1.390738986	Down	1.27E-119	6.96E-119	AAEL003933-PA OS=Aedes aegypti GN=DBLOX PE=4 SV=2
CCG017480.1	3123	29	789	0.39901451	8.831918147	4.468213699	Up	6.66E-13	1.69E-12	AAEL014084-PA OS=Aedes aegypti GN=AAEL014084 PE=3 SV=1
CCG009650.1	1150	261	789	9.75226151	23.98441772	1.298288697	Up	8.02E-14	2.58E-13	AAEL014548-PA (Fragment) OS=Aedes aegypti GN=TPX3 PE=4 SV=2
CCG022000.1	604	3128	789	222.532003	45.66569598	-2.284830103	Down	0	0	AAEL005538-PA OS=Aedes aegypti GN=AAEL005538 PE=4 SV=1
CCG017153.1	1218	155	790	5.46823388	22.67408756	2.051897642	Up	3.42E-13	9.27E-13	AAEL012705-PA OS=Aedes aegypti GN=AAEL012705 PE=4 SV=1
CCG011986.1	4266	162	790	1.63176207	6.473754958	1.988172044	Up	0	0	Condensin complex subunit 1 OS=Aedes aegypti GN=AAEL004634 PE=3 SV=1
CCG023380.1	1521	254	790	7.17574792	18.15715888	1.33933736	Up	2.35E-13	6.67E-13	AAEL011123-PA OS=Aedes aegypti GN=AAEL011123 PE=3 SV=1
CCG017471.1	4897	270	790	2.36917058	5.639583143	1.25120645	Up	2.59E-13	7.26E-13	AAEL010940-PA (Fragment) OS=Aedes aegypti GN=AAEL010940 PE=4 SV=1
CCG017708.1	1161	29	791	1.07331809	23.81739615	4.471866093	Up	6.66E-13	1.69E-12	AAEL005819-PA OS=Aedes aegypti GN=AAEL005819 PE=4 SV=1
CCG003752.1	7566	266	791	1.5106991	3.654770939	1.274564653	Up	4.09E-14	1.40E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006312 PE=4 SV=1
CCG008415.1	1902	30	792	0.67775607	14.55675878	4.424779228	Up	3.71E-13	9.93E-13	AAEL003088-PA OS=Aedes aegypti GN=AAEL003088 PE=4 SV=1
CCG007415.1	2112	1864	792	37.9240461	13.10935379	-1.532516321	Down	4.69E-153	2.67E-152	AAEL011574-PA OS=Aedes aegypti GN=AAEL011574 PE=4 SV=1
CCG013980.1	903	51	793	2.42686209	30.69979344	3.661064918	Up	0	0	AAEL010412-PA OS=Aedes aegypti GN=AAEL010412 PE=4 SV=1

CCG017796.1	1038	98	793	4.05687282	26.70704574	2.718780415	Up	0	0	Nuclear receptor FTZ-F1 protein OS=Aedes aegypti GN=FTZ-F1 PE=2 SV=1
CCG023339.1	987	294	793	12.7994954	28.08704507	1.133817915	Up	0	0	AAEL005390-PA (Fragment) OS=Aedes aegypti GN=AAEL005390 PE=4 SV=1
CCG009775.1	2475	3901	793	67.7272464	11.20077312	-2.596138024	Down	0	0	AAEL006749-PA OS=Aedes aegypti GN=AAEL006749 PE=4 SV=1
CCG021295.1	1965	68	794	1.48699336	14.12563448	3.24784556	Up	0	0	AAEL007692-PA OS=Aedes aegypti GN=AAEL007692 PE=4 SV=1
CCG025320.1	1003	101	794	4.32696231	27.67385021	2.677096918	Up	0	0	AAEL007038-PA OS=Aedes aegypti GN=AAEL007038 PE=4 SV=1
CCG009360.1	1524	119	794	3.35524831	18.21317044	2.440490638	Up	0	0	AAEL008133-PA OS=Aedes aegypti GN=AAEL008133 PE=4 SV=1
CCG016736.1	1642	147	794	3.84686418	16.90430679	2.135636056	Up	7.02E-14	2.29E-13	AAEL011059-PA OS=Aedes aegypti GN=AAEL011059 PE=4 SV=1
CCG027417.1	797	284	794	15.3116746	34.82668978	1.185561281	Up	0	0	AAEL014368-PA OS=Aedes aegypti GN=AAEL014368 PE=4 SV=1
CCG022628.1	4077	290	794	3.05646874	6.808160843	1.155399311	Up	1.62E-13	4.77E-13	AAEL001140-PA (Fragment) OS=Aedes aegypti GN=AAEL001140 PE=3 SV=1
CCG027962.1	1359	142	795	4.48984718	20.45020606	2.187377134	Up	2.62E-13	7.35E-13	AAEL007277-PA (Fragment) OS=Aedes aegypti GN=AAEL007277 PE=4 SV=1
CCG016763.1	1446	258	796	7.66679912	19.24397532	1.327710569	Up	1.69E-13	4.97E-13	-
CCG004193.1	1197	2191	796	78.6522043	23.24710803	-1.758435945	Down	8.65E-219	5.22E-218	Acetyl-CoA acetyltransferase OS=Aedes aegypti PE=2 SV=1
CCG011597.1	845	69	797	3.50877123	32.97248117	3.232224661	Up	0	0	AAEL002108-PA OS=Aedes aegypti GN=AAEL002108 PE=4 SV=1
CCG028505.1	3889	86	797	0.95021784	7.16424443	2.914484363	Up	0	0	DNA polymerase subunit gamma 1, mitochondrial OS=Culex quinquefasciatus GN=CpipJ_CPIJ012266 PE=3 SV=1
CCG028617.1	2037	127	797	2.67901635	13.67783338	2.352064431	Up	1.57E-13	4.66E-13	AAEL001455-PA OS=Aedes aegypti GN=AAEL001455 PE=4 SV=1

CCG004368.1	702	147	797	8.99793588	39.6890977	2.141076773	Up	7.02E-14	2.29E-13	AAEL009116-PA OS=Aedes aegypti GN=AAEL009116 PE=4 SV=1
CCG005524.1	2589	164	797	2.72191444	10.76158617	1.983197113	Up	1.14E-13	3.54E-13	AAEL000922-PA OS=Aedes aegypti GN=AAEL000922 PE=4 SV=1
CCG016703.1	1264	172	797	5.84714743	22.04252103	1.914484363	Up	1.35E-13	4.09E-13	AAEL009222-PA OS=Aedes aegypti GN=AAEL009222 PE=4 SV=1
CCG001049.1	1362	0	798	0.001	20.48216216	14.3220804	Up	4.44E-16	1.64E-15	AAEL009018-PA OS=Aedes aegypti GN=AAEL009018 PE=3 SV=1
CCG005672.2	1698	137	798	3.46693383	16.42915481	2.244526057	Up	2.90E-13	8.04E-13	AAEL011982-PA OS=Aedes aegypti GN=AAEL011982 PE=4 SV=1
CCG002244.1	1614	237	798	6.30968222	17.28420376	1.453814891	Up	2.32E-13	6.62E-13	AAEL004530-PA OS=Aedes aegypti GN=AAEL004530 PE=4 SV=1
CCG020733.1	6117	1513	798	10.6282832	4.560520658	-1.220638132	Down	2.63E-88	1.36E-87	AAEL000656-PA OS=Aedes aegypti GN=AAEL000656 PE=4 SV=1
CCG013072.1	845	31	799	1.57640447	33.05522265	4.390168586	Up	2.40E-13	6.77E-13	AAEL009836-PA OS=Aedes aegypti GN=AAEL009836 PE=4 SV=1
CCG016293.1	1428	73	799	2.19663209	19.55998819	3.154540338	Up	0	0	AAEL001111-PA OS=Aedes aegypti GN=AAEL001111 PE=4 SV=1
CCG003659.1	1776	143	799	3.45983787	15.7272878	2.18449356	Up	0	0	AAEL008947-PA (Fragment) OS=Aedes aegypti GN=AAEL008947 PE=4 SV=1
CCG027168.1	1607	198	799	5.29434191	17.38124651	1.715008277	Up	0	0	AAEL013902-PA OS=Aedes aegypti GN=AAEL013902 PE=4 SV=1
CCG018391.1	777	2	800	0.11060421	35.99307776	8.346169394	Up	9.36E-05	0.00015349	-
CCG024391.1	1128	131	800	4.99027946	24.79310409	2.312746392	Up	1.26E-13	3.87E-13	AAEL003385-PA OS=Aedes aegypti GN=AAEL003385 PE=4 SV=1
CCG003609.1	897	133	800	6.37120926	31.1779503	2.290886958	Up	9.90E-14	3.11E-13	AAEL012673-PA OS=Aedes aegypti GN=AAEL012673 PE=4 SV=1
CCG017712.1	1875	206	800	4.72094151	14.91553142	1.659668866	Up	1.10E-13	3.44E-13	AAEL005140-PA OS=Aedes aegypti GN=AAEL005140 PE=4 SV=1
CCG004601.1	1536	37	801	1.03507824	18.23019511	4.138518271	Up	5.17E-14	1.74E-13	AAEL010501-PA OS=Aedes aegypti GN=AAEL010501 PE=4 SV=1

CCG007442.1	855	63	801	3.16619097	32.75038561	3.370691713	Up	0	0	male-specific transcription factor FRU-MA OS=Culex quinquefasciatus GN=CpipJ_CPIJ010851 PE=4 SV=1
CCG022621.1	3135	160	801	2.19303271	8.931923348	2.026043541	Up	4.24E-14	1.45E-13	AAEL005806-PA (Fragment) OS=Aedes aegypti GN=AAEL005806 PE=4 SV=1
CCG000601.1	1074	178	801	7.12161337	26.07223435	1.872238205	Up	1.15E-13	3.57E-13	AAEL001449-PA OS=Aedes aegypti GN=AAEL001449 PE=4 SV=1
CCG015974.1	1221	1582	801	55.6741361	22.9333167	-1.279562248	Down	3.05E-99	1.61E-98	AGAP003014-PA OS=Anopheles gambiae GN=AgaP_AGAP003014 PE=4 SV=4
CCG017826.1	576	1892	801	141.143642	48.61385364	-1.537724737	Down	4.23E-156	2.42E-155	AAEL009257-PA OS=Aedes aegypti GN=AAEL009257 PE=4 SV=1
CCG023568.1	2619	30	802	0.49220773	10.70505459	4.442881035	Up	3.71E-13	9.94E-13	AAEL013025-PA OS=Aedes aegypti GN=AAEL013025 PE=4 SV=1
CCG002104.1	2616	55	802	0.90341567	10.71733103	3.568411917	Up	0	0	AAEL003153-PA OS=Aedes aegypti GN=AAEL003153 PE=4 SV=1
CCG025956.1	3430	281	802	3.52026106	8.173917776	1.21534531	Up	0	0	Na-K-Cl cotransporter OS=Aedes aegypti PE=2 SV=1
CCG011231.2	1458	69	803	2.03354711	19.25342678	3.243044925	Up	0	0	AAEL015034-PA OS=Aedes aegypti GN=AAEL015034 PE=4 SV=1
CCG010441.1	1287	92	803	3.07165158	21.8115744	2.828007425	Up	1.44E-14	5.12E-14	AAEL007594-PA OS=Aedes aegypti GN=AAEL007594 PE=3 SV=1
CCG014371.1	769	305	803	17.0426126	36.50389629	1.098903949	Up	5.30E-13	1.37E-12	AAEL011737-PA OS=Aedes aegypti GN=AAEL011737 PE=4 SV=1
CCG021478.1	741	87	804	5.04502957	37.93043796	2.910421399	Up	0	0	AAEL012372-PA OS=Aedes aegypti GN=AAEL012372 PE=4 SV=1
CCG016699.1	1020	226	804	9.52074512	27.55534757	1.533185933	Up	2.04E-14	7.16E-14	AAEL009225-PA OS=Aedes aegypti GN=AAEL009225 PE=4 SV=1
CCG005882.1	1119	1929	804	74.0738321	25.117475	-1.560272533	Down	1.53E-162	8.80E-162	AAEL012098-PA (Fragment) OS=Aedes aegypti GN=AAEL012098 PE=4 SV=1

CCG015923.1	2478	91	805	1.5779846	11.35650234	2.847363537	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005532 PE=4 SV=1
CCG011701.1	1116	124	805	4.77441496	25.21631971	2.400961866	Up	1.61E-13	4.77E-13	Transcription factor 11B OS=Culex quinquefasciatus GN=CpipJ_CPIJ011738 PE=4 SV=1
CCG019519.1	1950	170	805	3.74607943	14.43149374	1.945767241	Up	3.53E-13	9.52E-13	AAEL008037-PA OS=Aedes aegypti GN=AAEL008037 PE=3 SV=1
CCG015871.1	474	2703	805	245.036271	59.3700692	-2.04518762	Down	0	0	AAEL007135-PA OS=Aedes aegypti GN=AAEL007135 PE=4 SV=1
CCG023685.1	2163	2854	805	56.6970054	13.01036191	-2.123611443	Down	0	0	AAEL012552-PA OS=Aedes aegypti GN=AAEL012552 PE=4 SV=1
CCG018163.1	2730	53	806	0.83421096	10.32101505	3.629028778	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009453 PE=4 SV=1
CCG017010.1	660	135	806	8.7892639	42.69147133	2.280133635	Up	1.90E-13	5.48E-13	Probable ER retained protein OS=Aedes aegypti PE=2 SV=1
CCG013478.1	1474	172	806	5.01410743	19.11558418	1.930684478	Up	1.35E-13	4.09E-13	AAEL014705-PA OS=Aedes aegypti GN=AAEL014705 PE=4 SV=1
CCG010283.1	4610	1994	806	18.5860414	6.11201108	-1.604500462	Down	6.38E-175	3.72E-174	AAEL002683-PA OS=Aedes aegypti GN=AAEL002683 PE=4 SV=1
CCG019008.1	2196	84	807	1.64365105	12.84668914	2.966420644	Up	0	0	AAEL004387-PA OS=Aedes aegypti GN=AAEL004387 PE=4 SV=1
CCG009339.1	1719	187	807	4.67442721	16.41147723	1.811843607	Up	3.31E-13	9.04E-13	AAEL003101-PA OS=Aedes aegypti GN=AAEL003101 PE=4 SV=1
CCG025934.1	822	3524	807	184.215748	34.32035201	-2.424260142	Down	0	0	AAEL004081-PA OS=Aedes aegypti GN=AAEL004081 PE=4 SV=1
CCG000380.1	1176	2	808	0.07307778	24.01895207	8.360524687	Up	9.36E-05	0.00015339	AAEL005606-PA OS=Aedes aegypti GN=AAEL005606 PE=4 SV=1
CCG026773.1	870	82	808	4.05002096	32.46699728	3.002972682	Up	0	0	AAEL010747-PA OS=Aedes aegypti GN=AAEL010747 PE=4 SV=1
CCG022360.1	786	83	808	4.5375165	35.93675271	2.985485255	Up	1.32E-13	4.02E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009358 PE=4 SV=1

CCG002018.1	2009	114	808	2.43830251	14.05987438	2.527634672	Up	8.48E-14	2.70E-13	AAEL010816-PA OS=Aedes aegypti GN=AAEL010816 PE=3 SV=1
CCG025936.1	1625	1342	808	35.4863901	17.38233085	-1.02964427	Down	5.09E-60	2.45E-59	AAEL006186-PA OS=Aedes aegypti GN=AAEL006186 PE=4 SV=1
CCG010040.1	627	71	809	4.86579132	45.10565536	3.212561977	Up	0	0	AAEL000604-PA OS=Aedes aegypti GN=AAEL000604 PE=4 SV=1
CCG025664.1	1341	176	809	5.6395774	21.08966884	1.902877478	Up	0	0	AAEL012081-PA OS=Aedes aegypti GN=AAEL012081 PE=4 SV=1
CCG016967.1	1292	136	810	4.52312996	21.91656671	2.27662846	Up	1.70E-13	4.98E-13	AAEL006788-PA (Fragment) OS=Aedes aegypti GN=AAEL006788 PE=2 SV=1
CCG009692.1	1632	193	810	5.08159239	17.35061531	1.771634264	Up	6.08E-14	2.02E-13	AAEL001705-PA OS=Aedes aegypti GN=AAEL001705 PE=4 SV=1
CCG026762.1	1275	194	810	6.53814001	22.2087876	1.764178459	Up	5.97E-13	1.53E-12	AAEL005717-PA OS=Aedes aegypti GN=AAEL005717 PE=4 SV=1
CCG028572.1	1166	207	810	7.62841772	24.28490925	1.670604344	Up	6.88E-15	2.47E-14	AAEL006458-PA OS=Aedes aegypti GN=AAEL006458 PE=3 SV=1
CCG009950.1	1044	306	810	12.5945774	27.12280094	1.106703459	Up	0	0	AAEL000055-PA OS=Aedes aegypti GN=AAEL000055 PE=4 SV=1
CCG024565.2	3003	326	810	4.66471311	9.429305423	1.015363147	Up	6.17E-13	1.58E-12	Aldehyde dehydrogenase OS=Aedes aegypti GN=AAEL012162 PE=3 SV=1
CCG023261.1	297	2810	810	406.548668	95.34075483	-2.092263114	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002897 PE=3 SV=1
CCG001922.1	3958	99	811	1.07478619	7.163002138	2.736514688	Up	0	0	AAEL001554-PA OS=Aedes aegypti GN=AAEL001554 PE=4 SV=1
CCG014483.1	429	139	811	13.9225947	66.08662579	2.246930235	Up	1.46E-13	4.37E-13	AAEL012955-PA OS=Aedes aegypti GN=AAEL012955 PE=4 SV=1
CCG003849.1	27472	2583	811	4.04014358	1.032002128	-1.96896062	Down	9.64E-301	6.09E-300	AAEL012764-PA OS=Aedes aegypti GN=AAEL012764 PE=3 SV=1
CCG016614.1	1318	98	813	3.1950182	21.56379288	2.754714902	Up	0	0	AAEL006033-PA OS=Aedes aegypti GN=AAEL006033 PE=4 SV=1
CCG010343.1	1133	201	813	7.62305089	25.08480054	1.718373055	Up	0	0	AAEL003844-PA OS=Aedes aegypti GN=GALE5 PE=4 SV=1

CCG007156.1	1743	31	814	0.7642351	16.32589632	4.417001878	Up	2.40E-13	6.77E-13	AAEL011085-PA OS=Aedes aegypti GN=AAEL011085 PE=4 SV=1
CCG000585.1	915	134	814	6.29283545	31.09949431	2.305108998	Up	0	0	AAEL006516-PA OS=Aedes aegypti GN=AAEL006516 PE=4 SV=1
CCG019430.1	1395	222	814	6.83819433	20.39859304	1.576782322	Up	0	0	AAEL005395-PA OS=Aedes aegypti GN=AAEL005395 PE=4 SV=1
CCG005682.1	384	292	814	32.6749024	74.10426378	1.181373629	Up	0	0	-
CCG025255.1	1521	62	815	1.75156052	18.73175251	3.418773143	Up	0	0	AAEL007468-PA OS=Aedes aegypti GN=AAEL007468 PE=4 SV=1
CCG006888.1	2037	64	815	1.35005548	13.98674304	3.372969453	Up	0	0	AAEL001455-PA OS=Aedes aegypti GN=AAEL001455 PE=4 SV=1
CCG020185.1	2103	287	815	5.86415304	13.54778677	1.208062526	Up	2.87E-13	7.98E-13	AAEL001331-PA OS=Aedes aegypti GN=AAEL001331 PE=2 SV=1
CCG018554.3	1450	9227	815	273.435684	19.64896246	-3.798676488	Down	0	0	Hypothetical conserved protein OS=Glossina morsitans morsitans PE=2 SV=1
CCG024897.1	1353	168	816	5.33548811	21.083484	1.982421123	Up	1.45E-13	4.33E-13	AAEL010491-PB OS=Aedes aegypti GN=AAEL010491 PE=4 SV=1
CCG003824.3	1773	291	816	7.0525622	16.08908846	1.189863203	Up	6.00E-14	2.00E-13	AAEL001133-PA OS=Aedes aegypti GN=AAEL001133 PE=4 SV=1
CCG017793.1	30970	2307	816	3.20087755	0.921083431	-1.797063743	Down	2.36E-237	1.44E-236	AAEL002049-PB (Fragment) OS=Aedes aegypti GN=AAEL002049 PE=4 SV=1
CCG020758.1	738	0	817	0.001	38.70042293	15.24006171	Up	4.44E-16	1.65E-15	Serine 5-dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ016922 PE=3 SV=1
CCG022874.1	1711	219	817	5.49992512	16.69252608	1.601718412	Up	4.32E-13	1.14E-12	-
CCG025649.1	669	138	818	8.86371207	42.7442009	2.26974578	Up	1.49E-13	4.45E-13	AAEL002371-PA OS=Aedes aegypti GN=AAEL002371 PE=4 SV=1
CCG027418.1	1953	175	818	3.85033464	14.64202273	1.927059125	Up	3.55E-13	9.58E-13	AAEL012420-PA OS=Aedes aegypti GN=AAEL012420 PE=4 SV=1
CCG025225.1	1317	180	818	5.87285667	21.71288565	1.88641714	Up	3.32E-13	9.06E-13	AGAP013320-PA OS=Anopheles gambiae GN=AgaP_AGAP013320 PE=4 SV=1

CCG022589.1	5781	188	818	1.39738974	4.946526622	1.823681385	Up	1.80E-13	5.24E-13	AAEL002786-PA OS=Aedes aegypti GN=AAEL002786 PE=4 SV=1
CCG021037.1	1510	1611	818	45.8438692	18.93766252	-1.275470542	Down	1.58E-100	8.36E-100	AAEL008569-PA OS=Aedes aegypti GN=AAEL008569 PE=4 SV=1
CCG015170.1	1912	3220	818	72.365348	14.95599916	-2.274574736	Down	0	0	AAEL003434-PA OS=Aedes aegypti GN=AAEL003434 PE=4 SV=1
CCG014139.1	3978	95	819	1.02617516	7.197292277	2.810177237	Up	0	0	AGAP007129-PA OS=Anopheles gambiae GN=AGAP007129 PE=4 SV=4
CCG000707.2	1686	123	819	3.1348027	16.98151167	2.43751834	Up	0	0	AAEL003146-PA OS=Aedes aegypti GN=AAEL003146 PE=4 SV=1
CCG026674.1	501	309	819	26.5022914	57.14736263	1.108569818	Up	6.25E-13	1.59E-12	AAEL008924-PA OS=Aedes aegypti GN=AAEL008924 PE=4 SV=1
CCG006634.1	541	23	820	1.82680942	52.9866672	4.858231347	Up	4.01E-11	8.67E-11	AAEL008511-PA OS=Aedes aegypti GN=AAEL008511 PE=4 SV=1
CCG012058.1	2293	170	820	3.18571953	12.50143347	1.972402367	Up	3.53E-13	9.52E-13	AAEL001250-PA OS=Aedes aegypti GN=AAEL001250 PE=4 SV=1
CCG000209.1	1299	34	821	1.12468897	22.09449209	4.296088774	Up	8.82E-14	2.79E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009013 PE=4 SV=1
CCG019082.1	2191	181	821	3.549759	13.09938167	1.883705729	Up	1.53E-13	4.55E-13	AAEL014321-PA OS=Aedes aegypti GN=AAEL014321 PE=4 SV=1
CCG026620.1	2809	270	821	4.13023437	10.21742443	1.306736019	Up	2.59E-13	7.26E-13	AAEL004756-PA OS=Aedes aegypti GN=AAEL004756 PE=4 SV=1
CCG003485.1	687	277	821	17.3254971	41.77692173	1.26980945	Up	0	0	AAEL004071-PA OS=Aedes aegypti GN=AAEL004071 PE=4 SV=1
CCG028417.1	2235	1741	821	33.4721736	12.84149675	-1.382148872	Down	1.15E-122	6.30E-122	NADP+-dependent farnesol dehydrogenase 1 OS=Aedes aegypti GN=FOHSDR-1 PE=2 SV=1
CCG010052.1	1227	2800	821	98.0564441	23.39099041	-2.067659496	Down	0	0	AAEL003430-PA OS=Aedes aegypti GN=AAEL003430 PE=4 SV=1

CCG023139.1	2260	98	822	1.86328938	12.71491306	2.770597943	Up	0	0	Gastrin zinc finger protein XLCGF8.2DB OS=Culex quinquefasciatus GN=CpipJ_CPIJ012331 PE=4 SV=1
CCG028060.1	2392	129	822	2.31734773	12.01325397	2.374080532	Up	0	0	AAEL007488-PA OS=Aedes aegypti GN=AAEL007488 PE=4 SV=1
CCG018016.1	1132	204	822	7.74366242	25.38489709	1.712882446	Up	6.59E-14	2.17E-13	AAEL001366-PA OS=Aedes aegypti GN=AAEL001366 PE=4 SV=1
CCG011887.1	897	76	823	3.640691	32.07431637	3.139134311	Up	0	0	AAEL008621-PA OS=Aedes aegypti GN=AAEL008621 PE=4 SV=1
CCG018874.1	2930	316	823	4.63427855	9.819338493	1.083281076	Up	3.23E-13	8.83E-13	AAEL006207-PA OS=Aedes aegypti GN=AAEL006207 PE=4 SV=1
CCG016835.2	357	1846	823	222.190841	80.59008903	-1.463125013	Down	1.49E-141	8.40E-141	AAEL001864-PA OS=Aedes aegypti GN=4EBP PE=4 SV=1
CCG023190.1	4903	2578	823	22.5935092	5.867970994	-1.944974724	Down	2.85E-295	1.79E-294	SLC4-like anion exchanger OS=Aedes aegypti PE=2 SV=1
CCG016576.1	2442	58	824	1.02057519	11.79591321	3.530832736	Up	0	0	AAEL010080-PA OS=Aedes aegypti GN=AAEL010080 PE=4 SV=1
CCG021555.1	1815	66	824	1.5625358	15.87086505	3.344419612	Up	0	0	AAEL008320-PA OS=Aedes aegypti GN=AAEL008320 PE=4 SV=1
CCG002001.1	1972	77	824	1.67782432	14.6073124	3.12202719	Up	0	0	AAEL000385-PA OS=Aedes aegypti GN=AAEL000385 PE=4 SV=1
CCG025578.1	1425	149	824	4.49297576	20.21447022	2.169645211	Up	2.33E-13	6.63E-13	AAEL011214-PA OS=Aedes aegypti GN=AAEL011214 PE=4 SV=1
CCG003479.1	2226	83	825	1.60219585	12.95623465	3.015524082	Up	1.32E-13	4.01E-13	AAEL013014-PA OS=Aedes aegypti GN=AAEL013014 PE=4 SV=1
CCG004760.1	3268	1630	825	21.4322728	8.82514637	-1.280092736	Down	2.92E-102	1.55E-101	AAEL006436-PA OS=Aedes aegypti GN=AAEL006436 PE=4 SV=1
CCG004279.1	2739	3276	825	51.3942499	10.52960144	-2.287156129	Down	0	0	Integrin beta OS=Aedes aegypti GN=AAEL012466 PE=3 SV=1
CCG001913.1	1316	87	826	2.84070434	21.94189712	2.949367679	Up	0	0	AAEL004371-PA (Fragment) OS=Aedes aegypti GN=AAEL004371 PE=4 SV=1
CCG020944.1	1587	163	826	4.41340059	18.19504513	2.043583021	Up	1.80E-13	5.25E-13	AAEL001365-PA OS=Aedes aegypti GN=AAEL001365 PE=3 SV=1

CCG018682.1	372	200	826	23.1020079	77.62241025	1.748454985	Up	0	0	AAEL004835-PA OS=Aedes aegypti GN=AAEL004835 PE=4 SV=1
CCG004260.1	745	230	826	13.2658241	38.75910955	1.546821124	Up	3.85E-13	1.03E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018772 PE=4 SV=1
CCG025216.1	1062	270	826	10.9245088	27.18977082	1.315495578	Up	2.59E-13	7.27E-13	AAEL005825-PA OS=Aedes aegypti GN=AAEL005825 PE=2 SV=1
CCG016965.1	3864	2673	827	29.7251813	7.482012135	-1.990187399	Down	0	0	AAEL006791-PA OS=Aedes aegypti GN=AAEL006791 PE=4 SV=1
CCG016076.1	639	46	828	3.09328293	45.29804878	3.872238205	Up	3.40E-14	1.18E-13	AAEL000081-PA (Fragment) OS=Aedes aegypti GN=AAEL000081 PE=3 SV=1
CCG025507.1	644	187	828	12.4772366	44.94635585	1.848905701	Up	3.31E-13	9.04E-13	Salivary purine nucleosidase OS=Aedes albopictus PE=2 SV=1
CCG005989.1	1888	281	828	6.39538953	15.33127816	1.261373841	Up	0	0	AAEL009412-PA OS=Aedes aegypti GN=AAEL009412 PE=4 SV=1
CCG015544.1	1319	26	829	0.84701524	21.97150223	4.697101777	Up	4.60E-12	1.05E-11	AAEL000661-PA OS=Aedes aegypti GN=AAEL000661 PE=4 SV=1
CCG015138.1	1629	133	829	3.50827176	17.79030782	2.34225906	Up	9.90E-14	3.11E-13	C47D6 small nuclear ribonucleoprotein Prp31 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002179 PE=4 SV=1
CCG018462.1	46064	1799	829	1.67815545	0.629133628	-1.415437976	Down	2.54E-131	1.41E-130	AAEL009685-PA OS=Aedes aegypti GN=AAEL009685 PE=3 SV=1
CCG023164.1	786	66	830	3.60814565	36.91522865	3.354886611	Up	0	0	AAEL008772-PA OS=Aedes aegypti GN=AAEL008772 PE=4 SV=1
CCG005244.1	1285	194	830	6.48725954	22.58005426	1.799367888	Up	5.97E-13	1.53E-12	AAEL001945-PA OS=Aedes aegypti GN=AAEL001945 PE=4 SV=1
CCG023723.1	1657	49	831	1.27068014	17.53188171	3.786308026	Up	0	0	Pipsqueak protein OS=Drosophila melanogaster GN=psq PE=4 SV=1
CCG021435.1	2848	201	831	3.03262523	10.20025562	1.749966179	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016357 PE=4 SV=1
CCG008208.1	1266	269	831	9.13022007	22.9465466	1.329555508	Up	0	0	AAEL006663-PA OS=Aedes aegypti GN=AAEL006663 PE=4 SV=1

CCG002991.1	1636	132	832	3.4669957	17.77829234	2.358358803	Up	1.46E-13	4.36E-13	AAEL004385-PA OS=Aedes aegypti GN=AAEL004385 PE=4 SV=1
CCG012210.1	1068	136	832	5.47180141	27.23341411	2.315290081	Up	1.70E-13	4.98E-13	-
CCG025392.1	1191	232	832	8.37025897	24.42089528	1.544771927	Up	2.91E-13	8.06E-13	AAEL005410-PA OS=Aedes aegypti GN=AAEL005410 PE=4 SV=1
CCG002275.1	1782	51	833	1.22977355	16.34132691	3.732060547	Up	0	0	AAEL0006750-PA OS=Aedes aegypti GN=AAEL006750 PE=4 SV=1
CCG004556.1	2210	153	833	2.97482778	13.17658124	2.147098046	Up	0	0	AAEL011072-PA OS=Aedes aegypti GN=AAEL011072 PE=4 SV=1
CCG019560.1	1453	260	833	7.68900963	20.04146218	1.382118076	Up	3.22E-13	8.81E-13	AAEL010784-PA OS=Aedes aegypti GN=AAEL010784 PE=4 SV=1
CCG015801.1	570	284	833	21.4094818	51.08814834	1.25473877	Up	0	0	AAEL007494-PA OS=Aedes aegypti GN=AAEL007494 PE=4 SV=1
CCG001517.1	3534	71	834	0.86328556	8.249915911	3.256469658	Up	0	0	AAEL010401-PA OS=Aedes aegypti GN=AAEL010401 PE=4 SV=1
CCG027053.1	2231	211	834	4.0639238	13.0682218	1.685117589	Up	1.30E-13	3.98E-13	AAEL012291-PA OS=Aedes aegypti GN=AAEL012291 PE=4 SV=1
CCG005070.1	1995	221	834	4.76005581	14.61413676	1.618314218	Up	0	0	AAEL015141-PA OS=Aedes aegypti GN=AAEL015141 PE=4 SV=1
CCG017797.1	1876	337	834	7.71897685	15.54115289	1.009611996	Up	0	0	AAEL001480-PA OS=Aedes aegypti GN=AAEL001480 PE=4 SV=1
CCG005953.1	2124	115	835	2.32651576	13.7430137	2.56245554	Up	6.13E-14	2.03E-13	AAEL0006075-PA OS=Aedes aegypti GN=AAEL006075 PE=4 SV=1
CCG025311.1	873	149	835	7.33389514	33.43661066	2.188777071	Up	2.33E-13	6.62E-13	AAEL0006678-PA OS=Aedes aegypti GN=AAEL006678 PE=4 SV=1
CCG028260.1	1588	54	836	1.46118745	18.40372757	3.654784834	Up	0	0	AAEL000261-PA OS=Aedes aegypti GN=AAEL000261 PE=4 SV=1
CCG013693.1	2362	116	836	2.11028333	12.37303953	2.551691341	Up	1.16E-13	3.58E-13	AAEL012969-PA OS=Aedes aegypti GN=AAEL012969 PE=4 SV=1
CCG004573.1	4047	1791	836	19.0162576	7.221428066	-1.396877286	Down	3.59E-128	1.99E-127	Quick-to-court OS=Culex quinquefasciatus GN=CpipJ_CPIJ007694 PE=4 SV=1
CCG004252.1	2855	2107	836	31.7118147	10.23646914	-1.631302263	Down	3.41E-189	2.02E-188	AAEL012655-PA OS=Aedes aegypti GN=AAEL012655 PE=3 SV=1
CCG015513.1	542	0	837	0.001	53.98538313	15.72028122	Up	4.44E-16	1.64E-15	-

CCG017767.1	1992	91	837	1.96297482	14.68879401	2.903602376	Up	0	0	AAEL006227-PA OS=Aedes aegypti GN=AAEL006227 PE=4 SV=1
CCG000226.1	667	114	838	7.34415254	43.92059361	2.580229623	Up	8.48E-14	2.70E-13	AAEL007466-PA OS=Aedes aegypti GN=AAEL007466 PE=4 SV=1
CCG012471.1	4986	198	838	1.70637935	5.875458471	1.783763017	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_13311 PE=4 SV=1
CCG002968.1	885	240	838	11.6528094	33.10173552	1.506229042	Up	0	0	AAEL009700-PA OS=Aedes aegypti GN=AAEL009700 PE=4 SV=1
CCG017473.1	3174	256	838	3.46573789	9.229689961	1.413119638	Up	0	0	AAEL010928-PA (Fragment) OS=Aedes aegypti GN=AAEL010928 PE=4 SV=1
CCG006505.1	1293	168	839	5.58307457	22.68367688	2.022522781	Up	1.45E-13	4.33E-13	Ubiquitin fusion degradation protein 2 OS=Aedes aegypti PE=2 SV=1
CCG027531.2	2432	291	839	5.14152663	12.06003051	1.229964861	Up	6.00E-14	1.99E-13	AAEL012054-PA OS=Aedes aegypti GN=AAEL012054 PE=4 SV=1
CCG022090.1	2040	8769	839	184.706668	14.37744814	-3.68335641	Down	0	0	AAEL008513-PA (Fragment) OS=Aedes aegypti GN=AAEL008513 PE=4 SV=1
CCG005724.1	846	143	840	7.26320573	34.71034573	2.256687385	Up	0	0	AAEL002375-PA OS=Aedes aegypti GN=AAEL002375 PE=4 SV=1
CCG022131.1	1575	23	841	0.62749454	18.66661001	4.894713238	Up	4.01E-11	8.67E-11	Alkaline phosphatase OS=Aedes aegypti GN=AAEL003313 PE=3 SV=1
CCG001051.1	893	102	841	4.90807719	32.92263244	2.745849852	Up	0	0	AAEL002758-PA OS=Aedes aegypti GN=AAEL002758 PE=4 SV=1
CCG012434.1	1466	302	841	8.85188257	20.05450939	1.179870455	Up	4.59E-13	1.20E-12	AAEL000064-PA OS=Aedes aegypti GN=DCE PE=2 SV=1
CCG014229.1	2403	342	841	6.11554275	12.23466948	1.000422679	Up	1.09E-13	3.41E-13	AAEL003875-PA OS=Aedes aegypti GN=AAEL003875 PE=4 SV=1
CCG003750.1	921	41	843	1.91287635	31.99764095	4.06415002	Up	1.02E-13	3.19E-13	AAEL009650-PA OS=Aedes aegypti GN=AAEL009650 PE=4 SV=1
CCG022536.3	1904	252	843	5.68717076	15.47785048	1.444422101	Up	5.42E-14	1.81E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006732 PE=4 SV=1

CCG017645.1	1273	341	843	11.5103531	23.14990363	1.008074096	Up	2.98E-13	8.25E-13	AGAP012135-PA OS=Anopheles gambiae GN=AGAP012135 PE=4 SV=4
CCG017497.1	1215	2083	843	73.6674545	24.25500191	-1.602745099	Down	2.38E-182	1.40E-181	Short-chain dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ011768 PE=3 SV=1
CCG011449.1	1014	21	844	0.88990575	29.09742169	5.03109497	Up	1.76E-10	3.73E-10	AAEL000687-PA OS=Aedes aegypti GN=AAEL006887 PE=3 SV=1
CCG000562.1	1609	103	844	2.75070396	18.33734344	2.736911865	Up	0	0	Histone acetyltransferase Tipoo OS=Culex quinquefasciatus GN=CpipJ_CPIJ001206 PE=4 SV=1
CCG007719.1	723	191	844	11.3516173	40.80883208	1.845983564	Up	6.68E-14	2.19E-13	AAEL002532-PA OS=Aedes aegypti GN=Aael_AEL002532 PE=4 SV=1
CCG019585.1	4251	217	844	2.19346799	6.940669394	1.66186116	Up	1.47E-13	4.37E-13	AAEL001244-PA OS=Aedes aegypti GN=AAEL001244 PE=4 SV=1
CCG022110.1	2060	232	844	4.83930992	14.32271145	1.565431397	Up	2.91E-13	8.05E-13	AAEL012912-PA OS=Aedes aegypti GN=AAEL012912 PE=4 SV=1
CCG009108.1	2277	293	844	5.52926317	12.9577451	1.228655538	Up	4.19E-13	1.11E-12	AAEL014027-PA OS=Aedes aegypti GN=AAEL014027 PE=4 SV=1
CCG026016.1	1092	2365	844	93.0617421	27.01903443	-1.784212076	Down	7.50E-241	4.60E-240	AAEL011130-PA OS=Aedes aegypti GN=AAEL011130 PE=3 SV=1
CCG002877.1	1013	3169	844	134.423582	29.1261457	-2.206399552	Down	0	0	AAEL000377-PA OS=Aedes aegypti GN=AAEL006377 PE=4 SV=1
CCG012741.1	610	118	845	8.31217817	48.42580963	2.542477686	Up	0	0	AAEL005263-PA OS=Aedes aegypti GN=AAEL005263 PE=4 SV=1
CCG013855.1	3669	162	845	1.89727365	8.051170311	2.085270732	Up	0	0	Tetrapeptide repeat protein 37 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001407 PE=4 SV=1
CCG001750.1	3361	165	845	2.10949307	8.788974672	2.058798521	Up	1.31E-14	4.67E-14	AAEL001962-PA OS=Aedes aegypti GN=AAEL001962 PE=4 SV=1
CCG015171.1	2223	1581	845	30.5601216	13.28823386	-1.201500917	Down	6.06E-90	3.15E-89	AAEL003402-PA OS=Aedes aegypti GN=AAEL003402 PE=4 SV=1
CCG015746.1	1575	2215	845	60.4304522	18.75539293	-1.687970248	Down	9.46E-209	5.67E-208	Putative adenosine deaminase OS=Aedes albopictus PE=2 SV=1

CCG013889.1	851	75	846	3.78699189	34.75288149	3.198008366	Up	0	0	AAEL002108-PA OS=Aedes aegypti GN=AAEL002108 PE=4 SV=1
CCG020082.1	315	1386	846	189.066832	93.88794333	-1.009884485	Down	5.00E-60	2.41E-59	AAEL011156-PA OS=Aedes aegypti GN=AAEL011156 PE=4 SV=1
CCG019405.1	1218	3693	846	130.285082	24.28136465	-2.42375049	Down	0	0	Triosephosphate isomerase OS=Aedes aegypti GN=AAEL002542 PE=3 SV=1
CCG024960.1	3189	117	847	1.57650014	9.28493585	2.558166644	Up	9.44E-14	2.97E-13	AAEL007123-PA (Fragment) OS=Aedes aegypti GN=AAEL007123 PE=3 SV=2
CCG015569.1	2200	82	848	1.6015992	13.47482668	3.072681654	Up	0	0	AAEL008459-PA OS=Aedes aegypti GN=AAEL008459 PE=4 SV=1
CCG018251.1	2006	135	848	2.89178174	14.77797543	2.353418061	Up	1.90E-13	5.48E-13	AAEL007921-PA OS=Aedes aegypti GN=AAEL007921 PE=4 SV=1
CCG001736.1	1495	152	848	4.3688292	19.82917639	2.182306145	Up	8.39E-14	2.69E-13	Zinc-dependent metallopeptidase OS=Capsaspora owczarzaki (strain ATCC 30864) GN=CAOG_06679 PE=4 SV=1
CCG005418.1	1002	198	848	8.4910254	29.58544781	1.800877038	Up	0	0	AAEL010465-PA OS=Aedes aegypti GN=AAEL010465 PE=4 SV=1
CCG018513.1	1194	229	848	8.24126401	24.82798886	1.59102987	Up	2.46E-13	6.93E-13	AAEL006752-PA OS=Aedes aegypti GN=AAEL006752 PE=4 SV=1
CCG020277.1	1920	283	848	6.33355984	15.43990557	1.285575416	Up	0	0	AAEL008491-PA OS=Aedes aegypti GN=AAEL008491 PE=4 SV=1
CCG011353.1	736	2	849	0.11676558	40.3255122	8.431933947	Up	9.36E-05	0.00015357	-
CCG013411.1	1324	82	849	2.66126755	22.41659893	3.074381943	Up	0	0	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000594 PE=4 SV=1
CCG001912.1	1280	98	849	3.28987031	23.18716952	2.817224103	Up	0	0	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 SV=1
CCG015156.1	3018	244	849	3.47402758	9.834187203	1.50119661	Up	2.78E-13	7.73E-13	AAEL007624-PA OS=Aedes aegypti GN=REL2 PE=4 SV=1
CCG001009.1	3591	47	850	0.56239976	8.274724382	3.879043383	Up	0	0	terminal deoxycytidyl transferase rev1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007351 PE=4 SV=1

CCG019012.1	25109	6849	850	11.7208854	1.183421692	-3.30804541	Down	0	0	IP20063p OS=Drosophila melanogaster GN=CG33276-RB PE=4 SV=1
CCG021418.1	870	325	851	16.0519124	34.19482015	1.091032618	Up	1.19E-13	3.65E-13	AAEL010964-PA OS=Aedes aegypti GN=AAEL010964 PE=4 SV=1
CCG021911.1	7275	1466	851	8.65891834	4.089277462	-1.082340863	Down	7.69E-71	3.82E-70	Myosin-vIIra OS=Culex quinquefasciatus GN=CpipJ_CPIJ004181 PE=4 SV=1
CCG020775.1	843	2110	852	107.551768	35.33149681	-1.60600446	Down	3.13E-185	1.84E-184	AAEL012408-PA OS=Aedes aegypti GN=AAEL012408 PE=4 SV=1
CCG013446.1	3510	2843	852	34.8042608	8.485598806	-2.03617556	Down	0	0	AAEL005464-PA OS=Aedes aegypti GN=AAEL005464 PE=4 SV=1
CCG011917.1	1059	5097	852	206.814672	28.12507253	-2.878409814	Down	0	0	AAEL010260-PA OS=Aedes aegypti GN=AAEL010260 PE=4 SV=1
CCG014486.1	2553	51	853	0.85838483	11.68014496	3.766289793	Up	0	0	AAEL004200-PA OS=Aedes aegypti GN=AAEL004200 PE=3 SV=1
CCG024368.1	639	102	853	6.85901867	46.66574348	2.766289793	Up	0	0	AAEL011710-PA OS=Aedes aegypti GN=AAEL011710 PE=4 SV=1
CCG021213.1	483	330	853	29.3582038	61.73790908	1.072392921	Up	1.76E-13	5.16E-13	AAEL004813-PA OS=Aedes aegypti GN=AAEL004813 PE=4 SV=1
CCG007402.1	1013	4836	853	205.134883	29.43673256	-2.800883394	Down	0	0	AAEL004065-PA OS=Aedes aegypti GN=AAEL004065 PE=4 SV=1
CCG001999.1	399	20	854	2.15387141	74.82297835	5.118477369	Up	3.69E-10	7.70E-10	Short form D7 salivary protein SD7-1 OS=Aedes albopictus PE=2 SV=1
CCG021890.1	1635	8	856	0.21024947	18.30231493	6.44378019	Up	2.19E-06	3.96E-06	AAEL000053-PA OS=Aedes aegypti GN=AAEL000053 PE=4 SV=1
CCG020724.1	1566	182	856	4.99392829	19.10873877	1.93598555	Up	0	0	AAEL011251-PA OS=Aedes aegypti GN=AAEL011251 PE=4 SV=1
CCG024983.3	1875	192	856	4.40010082	15.95961862	1.858817689	Up	0	0	AAEL002349-PA OS=Aedes aegypti GN=AAEL002349 PE=4 SV=1
CCG021529.1	2271	223	856	4.2193971	13.17669966	1.64288029	Up	0	0	AAEL011269-PA OS=Aedes aegypti GN=AAEL011269 PE=4 SV=1
CCG021378.1	558	276	856	21.2538472	53.62775075	1.335255733	Up	0	0	AAEL008740-PA OS=Aedes aegypti GN=AAEL008740 PE=4 SV=1

CCG020400.2	5559	2125	857	16.4257395	5.389322395	-1.607782528	Down	7.83E-187	4.63E-186	AAEL003837-PA OS=Aedes aegypti GN=AAEL003837 PE=4 SV=1
CCG025258.1	1455	56	858	1.65381796	20.61457146	3.639792119	Up	0	0	AAEL007464-PA OS=Aedes aegypti GN=AAEL007464 PE=4 SV=1
CCG016249.1	1344	137	858	4.38009944	22.31711419	2.349114958	Up	2.90E-13	8.03E-13	AAEL003004-PA OS=Aedes aegypti GN=AAEL003004 PE=4 SV=1
CCG025189.1	567	308	858	23.3415842	52.89982623	1.180360501	Up	0	0	AAEL000950-PA OS=Aedes aegypti GN=AAEL000950 PE=4 SV=1
CCG021612.1	2132	130	859	2.62010577	14.08497174	2.426459712	Up	0	0	AAEL012932-PA OS=Aedes aegypti GN=AAEL012932 PE=4 SV=1
CCG001798.1	3414	142	860	1.78725903	8.806127131	2.300758934	Up	2.62E-13	7.34E-13	AAEL002465-PA (Fragment) OS=Aedes aegypti GN=AAEL002465 PE=4 SV=1
CCG012740.1	3168	1445	860	19.5995159	9.489936245	-1.046347724	Down	3.56E-66	1.74E-65	AAEL005763-PA (Fragment) OS=Aedes aegypti GN=AAEL005763 PE=4 SV=1
CCG011878.1	650	3154	860	208.502374	46.25248927	-2.172460891	Down	0	0	AAEL003593-PA OS=Aedes aegypti GN=AAEL003593 PE=4 SV=1
CCG001773.1	1536	78	861	2.18205684	19.5957528	3.166780412	Up	0	0	AAEL010805-PA OS=Aedes aegypti GN=AAEL010805 PE=4 SV=1
CCG027927.1	1788	107	861	2.57145504	16.83393529	2.710715645	Up	0	0	AGAP003794-PA OS=Anopheles gambiae GN=AgaP_AGAP003794 PE=4 SV=5
CCG003441.1	1503	140	861	4.00250356	20.02599887	2.322899614	Up	0	0	AAEL012851-PA OS=Aedes aegypti GN=AAEL012851 PE=4 SV=1
CCG010478.1	1386	36	862	1.116097	21.74172769	4.283932261	Up	6.13E-14	2.03E-13	AAEL008210-PA OS=Aedes aegypti GN=AAEL008210 PE=4 SV=1
CCG026557.1	900	46	862	2.19623088	33.48226064	3.930295307	Up	3.40E-14	1.18E-13	AAEL008316-PA OS=Aedes aegypti GN=AAEL008316 PE=4 SV=1
CCG023883.1	1554	5378	862	148.707357	19.39127064	-2.938996777	Down	0	0	AAEL001816-PA OS=Aedes aegypti GN=AAEL001816 PE=3 SV=1
CCG023670.1	1407	86	863	2.62643723	21.44207026	3.029265198	Up	0	0	AAEL002044-PA OS=Aedes aegypti GN=AAEL002044 PE=4 SV=1
CCG020559.1	2169	211	863	4.18008944	13.90917144	1.734430764	Up	1.30E-13	3.98E-13	AAEL009731-PA OS=Aedes aegypti GN=AAEL009731 PE=4 SV=1

CCG020519.1	1355	293	863	9.29161051	22.26493938	1.260773099	Up	4.19E-13	1.11E-12	AAEL008561-PA OS=Aedes aegypti GN=AAEL008561 PE=4 SV=1
CCG017599.1	655	1954	863	128.187575	46.05953108	-1.476684799	Down	8.66E-152	4.93E-151	Conserved insect protein (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG027580.1	1641	104	866	2.72324948	18.44842638	2.7600967	Up	0	0	AGAP004337-PA OS=Anopheles gambiae GN=AgaP_AGAP004337 PE=3 SV=5
CCG017180.1	2002	116	866	2.48975485	15.12181203	2.602555423	Up	1.16E-13	3.59E-13	AAEL013593-PA OS=Aedes aegypti GN=AAEL013593 PE=4 SV=1
CCG011504.1	1758	135	866	3.29972365	17.22063008	2.383720821	Up	1.90E-13	5.49E-13	AAEL007858-PA OS=Aedes aegypti GN=AAEL007858 PE=4 SV=1
CCG026942.1	901	177	866	8.44133521	33.6002971	1.992930868	Up	0	0	AAEL000055-PA OS=Aedes aegypti GN=AAEL000055 PE=4 SV=1
CCG008551.1	726	5230	866	309.547813	41.69954227	-2.892058813	Down	0	0	AAEL008289-PA OS=Aedes aegypti GN=AAEL008289 PE=4 SV=1
CCG009469.1	1665	9031	866	233.068873	18.18250311	-3.680133612	Down	0	0	AAEL008343-PA OS=Aedes aegypti GN=CYP4G35 PE=3 SV=1
CCG026309.1	2962	171	867	2.48069704	10.23255434	2.044348872	Up	3.42E-13	9.27E-13	AGAP010179-PA (Fragment) OS=Anopheles gambiae GN=AGAP010179 PE=4 SV=4
CCG005052.2	822	184	867	9.6185294	36.87205105	1.938639431	Up	3.33E-13	9.08E-13	AAEL002700-PA OS=Aedes aegypti GN=AAEL002700 PE=4 SV=1
CCG001445.1	1196	290	867	10.4190828	25.34182773	1.282292297	Up	1.62E-13	4.78E-13	AAEL003246-PA OS=Aedes aegypti GN=AAEL003246 PE=4 SV=1
CCG026619.1	1545	297	867	8.26020141	19.61736308	1.247882266	Up	2.44E-13	6.87E-13	AAEL004757-PA OS=Aedes aegypti GN=AAEL004757 PE=4 SV=1
CCG018249.1	1247	76	868	2.61884509	24.33342762	3.215936923	Up	0	0	AAEL013453-PA OS=Aedes aegypti GN=AAEL013453 PE=4 SV=1
CCG003614.1	1163	105	868	3.87946873	26.09095807	2.749618919	Up	0	0	AAEL006796-PA OS=Aedes aegypti GN=AAEL006796 PE=4 SV=1
CCG003048.1	2679	333	868	5.34114282	11.32653387	1.084486069	Up	4.41E-13	1.16E-12	AAEL010364-PA OS=Aedes aegypti GN=AAEL010364 PE=4 SV=1

CCG013611.1	1491	50	869	1.44097031	20.37474347	3.821669381	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005572 PE=4 SV=1
CCG017128.2	690	104	869	6.47659768	44.02716307	2.765085852	Up	0	0	AAEL008403-PA OS=Aedes aegypti GN=AAEL008403 PE=4 SV=1
CCG007617.1	1515	133	869	3.77226053	20.05197526	2.410243135	Up	9.90E-14	3.11E-13	AAEL013522-PA OS=Aedes aegypti GN=AAEL013522 PE=4 SV=1
CCG000839.1	1788	222	869	5.33516839	16.99034816	1.671109704	Up	0	0	Mi-primase inducer phosphatase 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001662 PE=4 SV=1
CCG012670.1	1818	224	869	5.29440074	16.70997938	1.658170649	Up	2.04E-13	5.87E-13	AAEL001398-PA (Fragment) OS=Aedes aegypti GN=AAEL001398 PE=4 SV=1
CCG027521.1	1329	238	869	7.69510672	22.85834651	1.570707807	Up	6.86E-13	1.73E-12	AAEL007985-PA OS=Aedes aegypti GN=AAEL007985 PE=4 SV=1
CCG012050.1	1440	8678	869	258.952331	21.09634897	-3.6176213	Down	0	0	AAEL000992-PA OS=Aedes aegypti GN=CYP6AG6 PE=3 SV=1
CCG001174.2	1758	194	870	4.74182509	17.3001711	1.867271952	Up	5.97E-13	1.53E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_10957 PE=4 SV=1
CCG011230.1	561	0	872	0.001	54.33799883	15.72967381	Up	4.44E-16	1.66E-15	5-hydroxyisourate hydrolase OS=Culex quinquefasciatus GN=CpipJ_CPIJ014057 PE=3 SV=1
CCG004581.1	729	3	872	0.17683018	41.81566165	7.885535028	Up	5.28E-05	8.85E-05	-
CCG003086.1	1360	83	872	2.62241763	22.41442452	3.095458097	Up	1.32E-13	4.02E-13	AAEL006235-PA OS=Aedes aegypti GN=AAEL006235 PE=4 SV=1
CCG000913.1	1387	193	872	5.97920604	21.9780947	1.878040491	Up	6.08E-14	2.02E-13	AAEL013048-PA OS=Aedes aegypti GN=AAEL013048 PE=4 SV=1
CCG010965.1	1453	200	872	5.9146228	20.97977794	1.826641339	Up	0	0	AAEL007971-PA OS=Aedes aegypti GN=AAEL007971 PE=4 SV=1
CCG016747.1	1491	224	872	6.45554698	20.44508206	1.663142607	Up	2.04E-13	5.88E-13	AAEL000559-PA OS=Aedes aegypti GN=AAEL000559 PE=4 SV=1
CCG008955.1	2931	321	872	4.70599959	10.40041533	1.144068041	Up	0	0	AAEL010200-PA OS=Aedes aegypti GN=AAEL010200 PE=4 SV=1

CCG011156.1	3552	134	873	1.62104292	8.591941335	2.406061857	Up	0	0	AAEL006614-PA OS=Aedes aegypti GN=AAEL006614 PE=4 SV=1
CCG027379.1	3784	337	873	3.82685004	8.065162691	1.075546266	Up	0	0	AAEL001252-PA OS=Aedes aegypti GN=AAEL001252 PE=4 SV=1
CCG026474.1	362	2	874	0.23740185	84.40202735	8.473802673	Up	9.36E-05	0.00015321	willstoni t1ZA OS=Drosopma willstoni GN=GK10379 PE=3 SV=1
CCG027570.1	2059	83	874	1.73214569	14.83901598	3.098763242	Up	1.32E-13	4.01E-13	AAEL002462-PA OS=Aedes aegypti GN=AAEL002462 PE=4 SV=1
CCG021275.1	1620	93	874	2.46678106	18.86020611	2.934643862	Up	4.04E-14	1.39E-13	AAEL010656-PA OS=Aedes aegypti GN=AAEL010656 PE=4 SV=1
CCG025132.1	1969	184	874	4.01545514	15.51728486	1.950240717	Up	3.33E-13	9.08E-13	AAEL014752-PA OS=Aedes aegypti GN=AAEL014752 PE=4 SV=1
CCG013554.1	1179	257	874	9.36660034	25.91478702	1.468178124	Up	4.14E-13	1.10E-12	AAEL000290-PA OS=Aedes aegypti GN=AAEL000290 PE=4 SV=1
CCG014281.1	1494	201	875	5.78106871	20.47422502	1.824400719	Up	0	0	AAEL007943-PA OS=Aedes aegypti GN=AAEL007943 PE=4 SV=1
CCG002121.1	1911	300	875	6.74564123	16.00653698	1.24663372	Up	1.10E-13	3.42E-13	AAEL001859-PA OS=Aedes aegypti GN=AAEL001859 PE=4 SV=1
CCG010484.1	897	1	876	0.04790383	34.13985558	9.477100263	Up	0.000587	0.00091638	AAEL013295-PA OS=Aedes aegypti GN=AAEL013295 PE=4 SV=1
CCG009343.1	1614	33	876	0.87856335	18.97363721	4.432706144	Up	1.31E-13	4.01E-13	AAEL003100-PA OS=Aedes aegypti GN=AAEL003100 PE=2 SV=1
CCG017891.1	1557	43	876	1.1867043	19.6682405	4.050835509	Up	8.46E-14	2.70E-13	AAEL000148-PA OS=Aedes aegypti GN=AAEL000148 PE=4 SV=1
CCG011760.1	1755	284	876	6.95350691	17.44925952	1.327353144	Up	0	0	AAEL003371-PA OS=Aedes aegypti GN=AAEL003371 PE=4 SV=1
CCG001005.1	651	317	876	20.9238185	47.0406305	1.168761233	Up	2.22E-13	6.35E-13	AAEL010486-PA OS=Aedes aegypti GN=AAEL010486 PE=4 SV=1
CCG019867.2	495	1624	876	140.975453	61.86555647	-1.188235654	Down	9.55E-91	4.97E-90	AAEL003423-PA OS=Aedes aegypti GN=AAEL003423 PE=4 SV=1
CCG007334.1	2102	142	877	2.90280795	14.58535144	2.328999117	Up	2.62E-13	7.35E-13	AAEL000077-PA OS=Aedes aegypti GN=AAEL000077 PE=4 SV=1
CCG002916.1	1878	215	877	4.91932531	16.32503127	1.730553387	Up	0	0	AAEL001848-PA (Fragment) OS=Aedes aegypti GN=AAEL001848 PE=4 SV=1

CCG016195.1	1801	1780	877	42.4686994	17.02299208	-1.31891529	Down	9.72E-117	5.29E-116	AAEL004957-PA OS=Aedes aegypti GN=AAEL004957 PE=4 SV=1
CCG023529.1	2306	34	878	0.63355203	13.31021986	4.392927492	Up	8.82E-14	2.80E-13	AAEL001635-PA OS=Aedes aegypti GN=AAEL001635 PE=4 SV=1
CCG024415.1	4326	80	878	0.79463217	7.095091772	3.158462238	Up	0	0	AAEL005539-PA OS=Aedes aegypti GN=AAEL005539 PE=4 SV=1
CCG003757.1	3396	352	878	4.4538712	9.038093936	1.020958715	Up	5.16E-13	1.34E-12	AAEL003445-PA OS=Aedes aegypti GN=AAEL003445 PE=4 SV=1
CCG023799.1	32813	1554	879	2.03501562	0.936468024	-1.119738229	Down	3.80E-79	1.92E-78	AAEL015040-PA OS=Aedes aegypti GN=AAEL015040 PE=4 SV=1
CCG022911.1	8415	1599	879	8.16501553	3.651613224	-1.160921665	Down	3.60E-86	1.85E-85	Protein translocase subunit secA OS=Harpegnathos saltator GN=EAI_00772 PE=4 SV=1
CCG011810.1	2499	206	880	3.54212298	12.31023752	1.79717239	Up	1.10E-13	3.44E-13	Mediator of RNA polymerase II transcription subunit 16 OS=Acromyrmex echinator GN=G5I_13705 PE=4 SV=1
CCG009038.1	4050	272	880	2.8858686	7.595872484	1.396210076	Up	5.36E-13	1.38E-12	AAEL002895-PA OS=Aedes aegypti GN=AAEL002895 PE=4 SV=1
CCG001769.1	1125	1946	880	74.328092	27.34514094	-1.442623077	Down	5.30E-146	3.00E-145	AAEL002077-PA OS=Aedes aegypti GN=AAEL002077 PE=4 SV=1
CCG023680.1	1122	29	881	1.11062594	27.4494134	4.627330418	Up	6.66E-13	1.69E-12	AAEL006407-PA OS=Aedes aegypti GN=AAEL006407 PE=4 SV=1
CCG024867.1	5550	82	881	0.63486815	5.549232763	3.127759408	Up	0	0	AAEL013080-PA OS=Aedes aegypti GN=AAEL013080 PE=4 SV=1
CCG005246.1	1344	119	882	3.80461192	22.94136913	2.592130286	Up	0	0	AAEL001917-PA OS=Aedes aegypti GN=AAEL001917 PE=4 SV=1
CCG001261.1	1325	1723	882	55.87687	23.27033971	-1.263758937	Down	7.31E-106	3.90E-105	AAEL004229-PA OS=Aedes aegypti GN=GSTT4 PE=3 SV=1
CCG006334.1	3957	112	883	1.21622701	7.800899265	2.681227909	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009355 PE=4 SV=1
CCG020315.1	2882	257	883	3.83179105	10.71067258	1.482958282	Up	4.14E-13	1.10E-12	AAEL004756-PA OS=Aedes aegypti GN=AAEL004756 PE=4 SV=1

CCG014421.1	1560	261	883	7.18916714	19.78728102	1.460676835	Up	8.02E-14	2.58E-13	AAEL014718-PA OS=Aedes aegypti GN=AAEL014718 PE=4 SV=1
CCG006078.1	1368	337	883	10.5853805	22.56444327	1.09197805	Up	0	0	AAEL007066-PA OS=Aedes aegypti GN=AAEL007066 PE=4 SV=1
CCG010744.1	1191	54	884	1.94824993	25.94720123	3.735328261	Up	0	0	AAEL006279-PA OS=Aedes aegypti GN=AAEL006279 PE=4 SV=1
CCG002681.1	1007	86	884	3.66970921	30.68829858	3.063951009	Up	0	0	AAEL006887-PA OS=Aedes aegypti GN=AAEL006887 PE=3 SV=1
CCG010680.1	1465	158	884	4.63427855	21.09427759	2.186435015	Up	0	0	AAEL002426-PA OS=Aedes aegypti GN=AAEL002426 PE=4 SV=1
CCG014948.1	1398	329	884	10.1123338	22.10523367	1.128271989	Up	0	0	AAEL001832-PA (Fragment) OS=Aedes aegypti GN=AAEL001832 PE=4 SV=1
CCG008622.1	3618	87	885	1.03326891	8.551153937	3.048903353	Up	0	0	AAEL007256-PA OS=Aedes aegypti GN=AAEL007256 PE=4 SV=1
CCG003171.1	1465	282	886	8.27130728	21.1420022	1.35392474	Up	5.45E-13	1.40E-12	Hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG014000.1	17550	2108	886	5.16126499	1.764845198	-1.548183059	Down	1.98E-175	1.16E-174	Calcium-activated potassium channel alpha chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ005769 PE=4 SV=1
CCG018419.1	1096	9138	886	358.264083	28.26006681	-3.664186635	Down	0	0	AAEL000445-PA OS=Aedes aegypti GN=AAEL000445 PE=4 SV=1
CCG007742.1	621	127	887	8.7876913	49.93235346	2.506418811	Up	1.57E-13	4.66E-13	-
CCG017398.1	753	225	887	12.8395621	41.17927158	1.681322307	Up	0	0	AAEL006560-PA OS=Aedes aegypti GN=AAEL006560 PE=3 SV=1
CCG025658.1	6309	295	887	2.00920458	4.914882152	1.290532354	Up	2.32E-13	6.61E-13	Pol-like protein OS=Biomphalaria glabrata PE=4 SV=1
CCG006378.1	1882	100	888	2.28319525	16.49465982	2.85287288	Up	0	0	AAEL002496-PA OS=Aedes aegypti GN=AAEL002496 PE=4 SV=1
CCG026130.1	1341	60	889	1.9225832	23.17517379	3.591462217	Up	0	0	AAEL002157-PA OS=Aedes aegypti GN=AAEL002157 PE=4 SV=1
CCG012837.1	1572	154	889	4.20950326	19.76966161	2.231566272	Up	1.82E-13	5.28E-13	AAEL003514-PA OS=Aedes aegypti GN=AAEL003514 PE=4 SV=1

CCG008027.1	354	312	889	37.8716305	87.79070071	1.212950594	Up	9.28E-14	2.94E-13	AAEL013700-PA OS=Aedes aegypti GN=AAEL013700 PE=4 SV=1
CCG020774.1	2529	109	890	1.85199726	12.30243825	2.731790405	Up	0	0	AAEL014046-PA OS=Aedes aegypti GN=AAEL014046 PE=4 SV=1
CCG018043.1	660	237	890	15.4300411	47.14070656	1.611231481	Up	2.32E-13	6.61E-13	AAEL012180-PA OS=Aedes aegypti GN=AAEL012180 PE=4 SV=1
CCG003735.1	1427	269	890	8.10011115	21.80298972	1.428512367	Up	0	0	AGAP003717-PA OS=Anopheles gambiae GN=AgaP_AGAP003717 PE=4 SV=3
CCG027132.1	7674	302	890	1.6910164	4.054321909	1.26156999	Up	4.59E-13	1.20E-12	AAEL000399-PA OS=Aedes aegypti GN=AAEL000399 PE=4 SV=1
CCG015571.1	798	143	891	7.70009029	39.03236166	2.341723489	Up	0	0	AAEL008461-PA OS=Aedes aegypti GN=AAEL008461 PE=2 SV=1
CCG005627.1	3431	145	891	1.81597538	9.078351677	2.321685735	Up	0	0	AGAP001132-PA OS=Anopheles gambiae GN=AgaP_AGAP001132 PE=4 SV=5
CCG003858.1	1569	145	891	3.97107171	19.85202333	2.321685735	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_12258 PE=3 SV=1
CCG018577.1	1780	275	891	6.63858259	17.49877787	1.398307017	Up	4.31E-13	1.14E-12	AAEL003140-PA OS=Aedes aegypti GN=AAEL003140 PE=4 SV=1
CCG007856.1	2688	54	892	0.86323128	11.60073768	3.748325602	Up	0	0	DNA-directed RNA polymerase OS=Aedes aegypti GN=AAEL013134 PE=3 SV=1
CCG023247.1	403	174	892	18.5526894	77.37663246	2.060269608	Up	1.78E-13	5.20E-13	-
CCG004753.1	621	182	892	12.5933844	50.21382106	1.995418464	Up	0	0	AAEL000206-PA OS=Aedes aegypti GN=AAEL000206 PE=4 SV=1
CCG009905.1	3516	238	892	2.90864529	8.868823345	1.60839534	Up	6.86E-13	1.73E-12	AAEL009048-PA OS=Aedes aegypti GN=AAEL009048 PE=4 SV=1
CCG005395.1	4128	257	892	2.67519908	7.553968721	1.497588555	Up	4.14E-13	1.10E-12	DNA-repair protein complementing XP-G cells OS=Culex quinquefasciatus GN=CpipJ_CPIJ004184 PE=4 SV=1
CCG001760.1	1131	258	892	9.80211453	27.57098398	1.491985848	Up	1.69E-13	4.97E-13	AAEL004520-PA OS=Aedes aegypti GN=AAEL004520 PE=4 SV=1

CCG015134.1	1612	287	892	7.65031876	19.34415811	1.338306177	Up	2.87E-13	7.98E-13	AAEL003537-PA OS=Aedes aegypti GN=AAEL003537 PE=4 SV=1
CCG006814.1	1050	75	893	3.06926676	29.73118205	3.276010878	Up	0	0	AAEL007077-PA OS=Aedes aegypti GN=AAEL007077 PE=4 SV=1
CCG025740.1	1023	144	893	6.04852569	30.51587601	2.334904567	Up	7.53E-14	2.44E-13	AAEL004285-PA OS=Aedes aegypti GN=AAEL004285 PE=4 SV=1
CCG000560.1	2509	276	893	4.72684207	12.44230417	1.396305112	Up	0	0	AAEL009415-PA OS=Aedes aegypti GN=AAEL009415 PE=4 SV=1
CCG021447.1	2105	311	893	6.34849761	14.83028083	1.224058799	Up	0	0	AAEL003528-PA OS=Aedes aegypti GN=AAEL003528 PE=4 SV=1
CCG023731.1	1131	316	894	12.0056907	27.63280233	1.202663477	Up	3.23E-13	8.83E-13	AAEL013363-PA OS=Aedes aegypti GN=AAEL013363 PE=4 SV=1
CCG007062.1	1881	53	895	1.21073681	16.6335235	3.780136621	Up	0	0	AAEL006544-PA OS=Aedes aegypti GN=AAEL006544 PE=4 SV=1
CCG025234.1	1309	145	895	4.75982545	23.90195394	2.328147986	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_07785 PE=4 SV=1
CCG022869.1	1636	2183	895	57.3367547	19.12448515	-1.584039339	Down	8.91E-188	5.27E-187	isocitrate dehydrogenase [NADP] OS=Culex quinquefasciatus GN=CpipJ_CPIJ003304 PE=3 SV=1
CCG028454.1	1587	3411	895	92.3564995	19.7149702	-2.227921964	Down	0	0	AAEL003419-PA OS=Aedes aegypti GN=AAEL003419 PE=4 SV=1
CCG022493.1	984	25	896	1.09171074	31.83192682	4.865811936	Up	9.32E-12	2.08E-11	Trypsin-like salivary secreted protein OS=Aedes albopictus PE=2 SV=1
CCG003709.1	1930	121	896	2.69395745	16.22933471	2.590804889	Up	0	0	Heat shock protein 70 BZ OS=Culex quinquefasciatus GN=CpipJ_CPIJ014283 PE=3 SV=1
CCG024639.1	2181	232	896	4.57082917	14.36158459	1.651687131	Up	2.91E-13	8.05E-13	AAEL011825-PA OS=Aedes aegypti GN=AAEL011825 PE=4 SV=1
CCG010481.1	2520	325	896	5.54173165	12.42960952	1.165372218	Up	1.19E-13	3.66E-13	AAEL013290-PA OS=Aedes aegypti GN=AAEL013290 PE=4 SV=1
CCG022588.1	1924	340	896	7.59340425	16.27994594	1.10027719	Up	8.86E-14	2.80E-13	AAEL013883-PA OS=Aedes aegypti GN=AAEL013883 PE=4 SV=1
CCG010279.1	1593	2003	896	54.0291139	19.66265913	-1.45827858	Down	1.15E-152	6.57E-152	Sterol carrier protein-X OS=Aedes aegypti PE=2 SV=1

CCG006752.1	2104	2031	896	41.4788645	14.8871749	-1.478306398	Down	5.99E-158	3.44E-157	AAEL006150-PA OS=Aedes aegypti GN=AAEL006150 PE=4 SV=1
CCG012242.1	4038	2334	896	24.8368897	7.75696285	-1.67892072	Down	3.54E-218	2.14E-217	AAEL002721-PA OS=Aedes aegypti GN=AAEL002721 PE=4 SV=1
CCG019648.1	2127	13	897	0.26262649	14.74263012	5.810837661	Up	6.34E-08	1.22E-07	Werner syndrome necase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007658 PE=4 SV=1
CCG014578.1	1566	66	897	1.81098498	20.02399378	3.466883259	Up	0	0	AAEL012611-PA OS=Aedes aegypti GN=AAEL012611 PE=4 SV=1
CCG001955.1	4338	270	897	2.67446481	7.228578669	1.434461782	Up	2.59E-13	7.27E-13	Dihydropyridine-sensitive r-type calcium channel OS=Culex quinquefasciatus GN=CpipJ_CPIJ001909 PE=4 SV=1
CCG019049.1	2201	261	898	5.09545694	14.26284986	1.484978842	Up	8.02E-14	2.58E-13	AAEL009147-PA OS=Aedes aegypti GN=AAEL009147 PE=4 SV=1
CCG001441.2	1371	1539	898	48.2351726	22.89754379	-1.074892678	Down	1.84E-73	9.23E-73	ATPase n2b OS=Culex quinquefasciatus GN=CpipJ_CPIJ009335 PE=4 SV=1
CCG009184.1	834	4176	898	215.157808	37.64092631	-2.515021158	Down	0	0	AAEL003235-PA OS=Aedes aegypti GN=AAEL003235 PE=4 SV=1
CCG002927.1	2268	105	899	1.98933957	13.85691835	2.800244992	Up	0	0	AAEL000305-PA OS=Aedes aegypti GN=AAEL000305 PE=4 SV=1
CCG004812.2	2111	129	899	2.62581514	14.88748973	2.503263254	Up	0	0	AAEL008839-PA OS=Aedes aegypti GN=AAEL008839 PE=4 SV=1
CCG013924.1	945	151	899	6.86606341	33.25660404	2.27608577	Up	0	0	AAEL000487-PA OS=Aedes aegypti GN=AAEL000487 PE=4 SV=1
CCG020700.1	2103	109	900	2.2271522	14.96074612	2.74791007	Up	0	0	AAEL003490-PA OS=Aedes aegypti GN=AAEL003490 PE=4 SV=1
CCG024497.1	1743	135	900	3.32812058	18.05074532	2.439278798	Up	1.90E-13	5.49E-13	AAEL004891-PA OS=Aedes aegypti GN=AAEL004891 PE=4 SV=1
CCG001407.1	1251	247	900	8.48403233	25.1498394	1.567727163	Up	0	0	AAEL002989-PA OS=Aedes aegypti GN=AAEL002989 PE=4 SV=1
CCG023848.1	1403	149	901	4.56342869	22.45004089	2.298527979	Up	2.33E-13	6.63E-13	AAEL013869-PA OS=Aedes aegypti GN=AAEL013869 PE=4 SV=1

CCG008770.1	2037	160	901	3.3751387	15.46264476	2.195768405	Up	4.24E-14	1.44E-13	AAEL010432-PA OS=Aedes aegypti GN=AAEL010432 PE=4 SV=1
CCG012550.1	726	178	901	10.5352793	43.38485864	2.041963069	Up	1.15E-13	3.58E-13	AAEL002774-PA OS=Aedes aegypti GN=AAEL002774 PE=4 SV=1
CCG000305.1	3267	316	901	4.15624002	9.641079698	1.213915751	Up	3.23E-13	8.83E-13	AAEL013386-PA OS=Aedes aegypti GN=AAEL013386 PE=4 SV=1
CCG015612.1	690	273	902	17.0010689	45.69908065	1.426539686	Up	1.53E-14	5.41E-14	AAEL012368-PA OS=Aedes aegypti GN=AAEL012368 PE=4 SV=1
CCG024783.1	1707	354	902	8.91112247	18.47238761	1.051691277	Up	0	0	AAEL010516-PA OS=Aedes aegypti GN=AAEL010516 PE=4 SV=1
CCG013101.1	3918	1518	902	16.6483045	8.04807699	-1.048659248	Down	1.14E-69	5.62E-69	AAEL005284-PA OS=Aedes aegypti GN=AAEL005284 PE=4 SV=1
CCG008547.1	2092	178	904	3.65612465	15.10625344	2.046758735	Up	1.15E-13	3.57E-13	AAEL000452-PA OS=Aedes aegypti GN=AAEL000452 PE=4 SV=1
CCG006293.1	1026	211	904	8.83685575	30.80144464	1.801392978	Up	1.30E-13	3.98E-13	AAEL011551-PA OS=Aedes aegypti GN=AAEL011551 PE=4 SV=1
CCG024870.1	1753	211	904	5.17205591	18.02754261	1.801392978	Up	1.30E-13	3.97E-13	AAEL004866-PA OS=Aedes aegypti GN=AAEL004866 PE=4 SV=1
CCG002084.1	2577	315	904	5.25241226	12.26320613	1.223284148	Up	4.28E-13	1.13E-12	AAEL001350-PA OS=Aedes aegypti GN=AAEL001350 PE=4 SV=1
CCG007780.1	627	1684	904	115.408346	50.40236396	-1.195184257	Down	6.81E-95	3.57E-94	AAEL003590-PA OS=Aedes aegypti GN=AAEL003590 PE=4 SV=1
CCG021483.1	2025	71	905	1.50659316	15.62332863	3.374340066	Up	0	0	AAEL013487-PA OS=Aedes aegypti GN=AAEL013487 PE=4 SV=1
CCG014521.1	1632	75	905	1.97471207	19.38556402	3.295268495	Up	0	0	AAEL007573-PA OS=Aedes aegypti GN=AAEL007573 PE=4 SV=1
CCG026502.1	4326	130	905	1.29127728	7.313277966	2.501719373	Up	0	0	DNA topoisomerase z-binding protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ010004 PE=4 SV=1
CCG002605.1	1514	3559	905	101.010096	20.89646003	-2.273169032	Down	0	0	AAEL014820-PA OS=Aedes aegypti GN=AAEL014820 PE=4 SV=1
CCG009319.3	1966	173	906	3.78116179	16.10996885	2.091052216	Up	3.60E-13	9.69E-13	AAEL010581-PA OS=Aedes aegypti GN=AAEL010581 PE=4 SV=1

CCG022055.1	1590	180	906	4.86449826	19.91962186	2.033827348	Up	3.32E-13	9.06E-13	Serine/threonine-protein phosphatase OS=Aedes aegypti GN=AAEL004040 PE=3 SV=1
CCG000364.1	1042	135	907	5.56709613	30.42913343	2.450456347	Up	1.90E-13	5.49E-13	AAEL009177-PA OS=Aedes aegypti GN=AAEL009177 PE=4 SV=1
CCG000925.1	1092	137	908	5.39089161	29.06787116	2.430829608	Up	2.90E-13	8.04E-13	AAEL013314-PA OS=Aedes aegypti GN=AAEL013314 PE=4 SV=1
CCG012806.1	1926	169	908	3.77044919	16.48084907	2.127982255	Up	0	0	AAEL007896-PA (Fragment) OS=Aedes aegypti GN=AAEL007896 PE=4 SV=1
CCG008922.1	2052	179	908	3.74833455	15.46886711	2.045045914	Up	0	0	AAEL000583-PA OS=Aedes aegypti GN=AAEL000583 PE=3 SV=1
CCG011080.1	1290	214	908	7.12831256	24.60629094	1.787394705	Up	0	0	AAEL013995-PA OS=Aedes aegypti GN=AAEL013995 PE=4 SV=1
CCG021486.1	1971	7245	908	157.948111	16.10457398	-3.293908283	Down	0	0	AAEL004680-PA OS=Aedes aegypti GN=AAEL004680 PE=4 SV=1
CCG004331.1	2304	22	909	0.41030129	13.79213263	5.071018069	Up	8.38E-11	1.80E-10	AAEL000116-PA OS=Aedes aegypti GN=AAEL000116 PE=4 SV=1
CCG028432.1	2387	100	909	1.80015646	13.31255701	2.886593498	Up	0	0	AAEL014827-PA OS=Aedes aegypti GN=AAEL014827 PE=4 SV=1
CCG025246.1	2903	179	909	2.64952893	10.94628784	2.046633911	Up	0	0	-
CCG008048.1	276	8182	909	1273.83467	115.1343246	-3.467788134	Down	0	0	AAEL001390-PA OS=Aedes aegypti GN=AAEL001390 PE=4 SV=1
CCG005728.1	3837	326	910	3.65080362	8.290860532	1.183307785	Up	6.17E-13	1.58E-12	AAEL002382-PA OS=Aedes aegypti GN=AAEL002382 PE=4 SV=1
CCG028063.1	1590	351	910	9.4857716	20.00756721	1.076708719	Up	4.44E-15	1.60E-14	AAEL000905-PA (Fragment) OS=Aedes aegypti GN=AAEL000905 PE=4 SV=2
CCG011877.1	582	4412	910	325.743074	54.65984856	-2.575181137	Down	0	0	AAEL003603-PA OS=Aedes aegypti GN=AAEL003603 PE=4 SV=1
CCG010865.1	2757	4820	910	75.1230036	11.5386405	-2.702781492	Down	0	0	AAEL012110-PA OS=Aedes aegypti GN=AAEL012110 PE=4 SV=1
CCG011754.1	1220	31	911	1.09185391	26.10409028	4.579424137	Up	2.40E-13	6.76E-13	AAEL008256-PA OS=Aedes aegypti GN=AAEL008256 PE=3 SV=1

CCG020318.1	1719	66	911	1.64979784	18.52646314	3.489226328	Up	0	0	AAEL004740-FA OS=Aedes aegypti GN=AAEL004740 PE=4 SV=1
CCG008962.1	3000	1944	911	27.844388	10.61566338	-1.391192056	Down	4.25E-138	2.38E-137	Heat shock protein 70 BZ OS=Culex quinquefasciatus GN=CpipJ_CPIJ014283 PE=3 SV=1
CCG010285.1	3346	184	912	2.36295014	9.528376693	2.011641262	Up	3.33E-13	9.08E-13	AAEL014497-FA OS=Aedes aegypti GN=AAEL014497 PE=4 SV=1
CCG012760.1	2835	219	912	3.31935516	11.24583718	1.760416158	Up	4.32E-13	1.14E-12	AAEL008333-FA OS=Aedes aegypti GN=AAEL008333 PE=4 SV=1
CCG023051.1	1912	248	912	5.57348022	16.67465921	1.581006908	Up	5.86E-14	1.95E-13	AAEL000214-FA OS=Aedes aegypti GN=AAEL000214 PE=4 SV=1
CCG013113.1	1023	342	912	14.3652485	31.16514997	1.117350703	Up	1.09E-13	3.41E-13	Putative arsenite-translocating ATPase OS=Glossina morsitans morsitans PE=2 SV=1
CCG006039.1	2822	361	912	5.49683706	11.29764295	1.039348191	Up	1.35E-13	4.08E-13	AAEL010837-FA OS=Aedes aegypti GN=AAEL010837 PE=4 SV=1
CCG018866.2	22677	1650	912	3.12651859	1.405915616	-1.153047091	Down	6.57E-88	3.40E-87	AAEL004669-FA OS=Aedes aegypti GN=AAEL004669 PE=4 SV=1
CCG015007.1	45084	1988	912	1.89477048	0.707167696	-1.421898824	Down	6.60E-146	3.73E-145	AAEL009685-FA OS=Aedes aegypti GN=AAEL009685 PE=3 SV=1
CCG028605.1	2016	189	913	4.02841262	15.83179895	1.97454183	Up	2.34E-13	6.64E-13	AAEL002006-FA OS=Aedes aegypti GN=AAEL002006 PE=4 SV=1
CCG019904.1	2352	198	913	3.61735011	13.57011339	1.907427634	Up	0	0	AAEL002386-FA OS=Aedes aegypti GN=AAEL002386 PE=4 SV=1
CCG004188.3	2211	1502	913	29.1906564	14.43550732	-1.015884844	Down	1.60E-65	7.83E-65	AAEL014217-FA OS=Aedes aegypti GN=AAEL014217 PE=4 SV=1
CCG010618.1	2493	136	914	2.34411709	12.81663256	2.450900718	Up	1.70E-13	4.98E-13	AAEL009919-FA OS=Aedes aegypti GN=AAEL009919 PE=4 SV=1
CCG014005.1	1317	187	914	6.10124554	24.26109717	1.991469099	Up	3.31E-13	9.03E-13	AAEL006617-FA OS=Aedes aegypti GN=AAEL006617 PE=4 SV=1
CCG011923.1	861	77	915	3.84282179	37.15078194	3.273154596	Up	0	0	Goodpasture antigen-binding protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000098 PE=4 SV=1
CCG005302.1	1230	120	915	4.19216923	26.00554736	2.633050541	Up	0	0	AAEL012705-FA OS=Aedes aegypti GN=AAEL012705 PE=4 SV=1

CCG003590.1	2004	87	916	1.86545255	15.9789329	3.098573496	Up	0	0	AAEL006227-PA OS=Aedes aegypti GN=AAEL006227 PE=4 SV=1
CCG010914.1	5808	119	916	0.88040606	5.513392136	2.646699229	Up	0	0	AAEL008377-PA OS=Aedes aegypti GN=AAEL008377 PE=4 SV=1
CCG024222.1	1521	83	917	2.34483101	21.07609454	3.168051696	Up	1.32E-13	4.02E-13	AAEL014161-PA OS=Aedes aegypti GN=AAEL014161 PE=4 SV=1
CCG000938.1	2353	93	917	1.6983363	13.62377382	3.003932316	Up	4.04E-14	1.39E-13	AAEL008572-PA OS=Aedes aegypti GN=AAEL008572 PE=4 SV=1
CCG006463.1	1865	134	917	3.08736967	17.18860043	2.477001937	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005688 PE=4 SV=1
CCG022394.1	1317	194	917	6.32963441	24.34072878	1.943178285	Up	5.97E-13	1.53E-12	AAEL007996-PA OS=Aedes aegypti GN=AAEL007996 PE=4 SV=1
CCG008300.1	2553	83	918	1.39697923	12.57019118	3.169624116	Up	1.32E-13	4.01E-13	AAEL006631-PA OS=Aedes aegypti GN=AAEL006631 PE=4 SV=1
CCG001277.1	2967	233	919	3.37443484	10.82799338	1.68204811	Up	0	0	AGAP010179-PA (Fragment) OS=Anopheles gambiae GN=AGAP010179 PE=4 SV=4
CCG027060.1	2685	144	920	2.30452208	11.97825498	2.377878253	Up	7.53E-14	2.44E-13	Anoctamin OS=Aedes aegypti GN=AAEL012806 PE=3 SV=1
CCG022772.1	1452	158	920	4.67577002	22.14987234	2.244022507	Up	0	0	AAEL003868-PA OS=Aedes aegypti GN=AAEL003868 PE=4 SV=1
CCG021113.1	1400	315	920	9.66819029	22.97258188	1.248595236	Up	4.28E-13	1.13E-12	AAEL009445-PA OS=Aedes aegypti GN=AAEL009445 PE=4 SV=1
CCG003850.1	2547	47	921	0.79292404	12.64097876	3.994781698	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015955 PE=4 SV=1
CCG022334.1	699	101	921	6.20878855	46.06090545	2.891159067	Up	0	0	AAEL011061-PA OS=Aedes aegypti GN=AAEL011061 PE=4 SV=1
CCG016430.1	648	158	921	10.4771884	49.6860693	2.245589802	Up	0	0	AAEL002375-PA OS=Aedes aegypti GN=AAEL002375 PE=4 SV=1
CCG005241.1	1119	177	921	6.79682129	28.77262994	2.081765	Up	0	0	AAEL002077-PA OS=Aedes aegypti GN=AAEL002077 PE=4 SV=1
CCG028498.1	2790	354	921	5.45207385	11.53999029	1.081765	Up	0	0	AAEL000569-PA OS=Aedes aegypti GN=AAEL000569 PE=4 SV=1

CCG024436.1	2797	211	922	3.24154952	11.52360786	1.829836956	Up	1.30E-13	3.98E-13	AAEL013624-PA OS=Aedes aegypti GN=AAEL013624 PE=4 SV=1
CCG004437.1	1135	269	922	10.1840164	28.39782483	1.479473782	Up	0	0	Splicing factor Uzar 38 kDa subunit OS=Acromyrmex echinator GN=G5I_01763 PE=4 SV=1
CCG024458.1	1755	361	922	8.83878871	18.36554483	1.055081117	Up	1.35E-13	4.08E-13	AAEL014698-PA OS=Aedes aegypti GN=AAEL014698 PE=4 SV=1
CCG006330.1	1416	132	923	4.00565323	22.78706883	2.508105922	Up	1.46E-13	4.36E-13	AAEL005074-PA OS=Aedes aegypti GN=AAEL005074 PE=4 SV=1
CCG025763.1	3165	158	923	2.1450926	10.1947834	2.248719293	Up	0	0	AAEL010501-PA OS=Aedes aegypti GN=AAEL010501 PE=4 SV=1
CCG014212.1	2378	208	923	3.75849655	13.56875082	1.852060323	Up	3.93E-13	1.05E-12	AAEL003463-PA OS=Aedes aegypti GN=AAEL003463 PE=4 SV=1
CCG009840.1	657	225	923	14.7156625	49.1118561	1.73871885	Up	0	0	AAEL010500-PA OS=Aedes aegypti GN=GST-2 PE=2 SV=1
CCG015208.1	1788	318	923	7.64226824	18.04613505	1.239617086	Up	1.20E-13	3.69E-13	AAEL015320-PA (Fragment) OS=Aedes aegypti GN=AAEL015320 PE=4 SV=1
CCG027346.1	588	37	924	2.70387786	54.93443493	4.34460888	Up	5.17E-14	1.74E-13	AAEL003950-PA OS=Aedes aegypti GN=AAEL003950 PE=4 SV=1
CCG022822.1	624	196	924	13.4969038	51.76514061	1.939352401	Up	7.93E-14	2.55E-13	Subolesin-like protein OS=Aedes albopictus PE=2 SV=1
CCG012500.1	2480	126	925	2.18313974	13.03887339	2.578342836	Up	1.34E-13	4.06E-13	Zinc finger protein z6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ016622 PE=4 SV=1
CCG024841.1	2699	197	925	3.13636077	11.98088404	1.93357094	Up	0	0	-
CCG014247.1	1538	224	925	6.25827084	21.0249714	1.748267837	Up	2.04E-13	5.88E-13	AAEL008550-PA OS=Aedes aegypti GN=AAEL008550 PE=4 SV=1
CCG024463.1	1672	245	925	6.2964025	19.33995575	1.61898482	Up	3.40E-13	9.24E-13	AAEL009149-PA OS=Aedes aegypti GN=AAEL009149 PE=4 SV=1
CCG010260.1	2712	58	926	0.91896925	11.93634377	3.699200592	Up	0	0	AAEL002918-PA OS=Aedes aegypti GN=AAEL002918 PE=4 SV=1
CCG001310.1	2730	169	926	2.66003119	11.8576426	2.156302151	Up	0	0	XPA-binding protein z OS=Culex quinquefasciatus GN=CpipJ_CPIJ005885 PE=4 SV=1

CCG026872.2	2541	261	926	4.41365633	12.73961601	1.529275591	Up	8.02E-14	2.58E-13	AGAP005366-PA OS=Anopheles gambiae GN=AGAP005366 PE=4 SV=3
CCG002735.1	2406	108	927	1.92881602	13.468962	2.80385123	Up	0	0	AAEL004526-PA OS=Aedes aegypti GN=AAEL004526 PE=4 SV=1
CCG009959.1	3180	190	927	2.56737408	10.19066747	1.988883124	Up	2.69E-13	7.52E-13	AAEL008656-PA OS=Aedes aegypti GN=AAEL008656 PE=4 SV=1
CCG000553.1	2200	326	927	6.3673334	14.73014662	1.210010578	Up	6.17E-13	1.58E-12	AAEL007307-PA OS=Aedes aegypti GN=AAEL007307 PE=4 SV=1
CCG002004.1	693	1793	927	111.175663	46.76237023	-1.24942104	Down	3.74E-108	2.01E-107	AAEL000235-PA OS=Aedes aegypti GN=AAEL000235 PE=4 SV=1
CCG025197.2	1659	172	928	4.45496947	19.55472022	2.134029444	Up	1.35E-13	4.09E-13	AAEL002341-PA OS=Aedes aegypti GN=AAEL002341 PE=4 SV=1
CCG024622.1	1692	223	928	5.66326881	19.17333383	1.759394299	Up	0	0	AAEL005120-PA OS=Aedes aegypti GN=AAEL005120 PE=4 SV=1
CCG009139.1	2085	252	928	5.19346433	15.55936731	1.583014275	Up	5.42E-14	1.81E-13	AAEL002798-PA OS=Aedes aegypti GN=AAEL002798 PE=4 SV=1
CCG027868.1	1290	363	928	12.0914835	25.14827972	1.056468461	Up	0	0	AAEL003248-PA OS=Aedes aegypti GN=AAEL003248 PE=4 SV=1
CCG024793.1	921	5939	928	277.087138	35.22397486	-2.975710118	Down	0	0	AAEL011088-PA OS=Aedes aegypti GN=AAEL011088 PE=4 SV=1
CCG014819.1	730	137	929	8.06418307	44.4879988	2.463815907	Up	2.90E-13	8.03E-13	AAEL003174-PA OS=Aedes aegypti GN=AAEL003174 PE=4 SV=1
CCG014479.1	1920	122	930	2.73036855	16.93291531	2.632662772	Up	8.04E-14	2.58E-13	Heat shock 70 Ab OS=Aedes aegypti GN=Hsp70Ab PE=3 SV=1
CCG022240.1	1018	205	930	8.65304086	31.93634322	1.88392001	Up	1.83E-13	5.31E-13	AAEL012389-PA OS=Aedes aegypti GN=AAEL012389 PE=4 SV=1
CCG016842.1	1359	304	930	9.6120672	23.92288256	1.315472596	Up	0	0	AAEL009636-PA OS=Aedes aegypti GN=AAEL009636 PE=4 SV=1
CCG021490.1	354	0	932	0.001	92.03704506	16.48992704	Up	4.44E-16	1.64E-15	AAEL010229-PA OS=Aedes aegypti GN=AAEL010229 PE=4 SV=1
CCG010419.1	981	302	932	13.2281956	33.2121447	1.328094609	Up	4.59E-13	1.20E-12	-
CCG021334.1	2157	117	933	2.33076446	15.12103488	2.697681755	Up	9.44E-14	2.98E-13	AAEL013400-PA OS=Aedes aegypti GN=AAEL013400 PE=3 SV=1

CCG023100.1	891	207	933	9.98286764	36.60614167	1.874559517	Up	6.88E-15	2.47E-14	DRE2 OS=Glossina morsitans morsitans PE=2 SV=1
CCG024734.1	2763	3	934	0.04665552	11.81723869	7.984629443	Up	5.28E-05	8.84E-05	AAEL003429-PA OS=Aedes aegypti GN=AAEL003429 PE=4 SV=1
CCG015367.1	1741	168	934	4.14641896	18.7541818	2.177274521	Up	1.45E-13	4.33E-13	AAEL006577-PA OS=Aedes aegypti GN=AAEL006577 PE=3 SV=1
CCG006733.1	1263	263	934	8.9477753	25.85196398	1.530672954	Up	1.40E-14	4.98E-14	AAEL006346-PA OS=Aedes aegypti GN=AAEL006346 PE=4 SV=1
CCG020695.1	1191	273	934	9.84948577	27.41480311	1.476834803	Up	1.53E-14	5.42E-14	Coractor required for Sp1 transcriptional activation subunit 9 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007281 PE=4 SV=1
CCG006074.1	2039	39	935	0.8218831	16.03040156	4.28573354	Up	7.33E-14	2.38E-13	AAEL007045-PA OS=Aedes aegypti GN=AAEL007045 PE=4 SV=1
CCG003483.1	833	189	935	9.74943558	39.23888209	2.008893334	Up	2.34E-13	6.64E-13	AAEL004067-PA OS=Aedes aegypti GN=AAEL004067 PE=4 SV=1
CCG001076.1	1432	220	935	6.60149554	22.82541116	1.789776045	Up	4.61E-13	1.20E-12	-
CCG020837.1	1263	163	936	5.54557937	25.9073215	2.223949769	Up	1.80E-13	5.25E-13	AAEL005191-PA OS=Aedes aegypti GN=AAEL005191 PE=4 SV=1
CCG013409.1	1044	212	936	8.72565492	31.34190331	1.844757469	Up	0	0	Insulin-like peptide 6 splice variant A OS=Aedes aegypti PE=2 SV=1
CCG006401.1	10126	2045	936	8.67796833	3.231379326	-1.425207204	Down	1.47E-150	8.36E-150	AAEL008485-PA OS=Aedes aegypti GN=AAEL008485 PE=4 SV=1
CCG004374.1	1966	83	937	1.8140834	16.66119295	3.19917901	Up	1.32E-13	4.02E-13	AAEL000429-PA OS=Aedes aegypti GN=AAEL000429 PE=4 SV=1
CCG008795.2	1717	148	937	3.70385598	19.07740555	2.364765076	Up	0	0	AAEL011816-PA OS=Aedes aegypti GN=AAEL011816 PE=4 SV=1
CCG019214.1	1875	277	937	6.34806213	17.46981618	1.460476275	Up	0	0	AAEL011778-PA OS=Aedes aegypti GN=AAEL011778 PE=4 SV=1
CCG021093.1	1875	102	938	2.33755356	17.48846059	2.903331974	Up	0	0	AAEL008400-PA OS=Aedes aegypti GN=AAEL008400 PE=4 SV=1
CCG012431.1	1947	362	938	7.98923674	16.84173786	1.075911429	Up	1.06E-12	2.62E-12	AAEL000070-PA OS=Aedes aegypti GN=AAEL000070 PE=4 SV=1
CCG008203.1	1209	6445	938	229.065293	27.12230241	-3.078207327	Down	0	0	AAEL012950-PA OS=Aedes aegypti GN=AAEL012950 PE=2 SV=1

CCG024289.1	1812	31	940	0.73513343	18.13508839	4.62463384	Up	2.40E-13	6.77E-13	AAEL004302-PA OS=Aedes aegypti GN=AAEL004562 PE=4 SV=1
CCG025047.1	1296	79	940	2.61929709	25.35554025	3.275049402	Up	0	0	AAEL013471-PA OS=Aedes aegypti GN=AAEL013471 PE=4 SV=1
CCG001190.1	2178	135	940	2.6634133	15.0875942	2.502014553	Up	1.90E-13	5.48E-13	AAEL004243-PA OS=Aedes aegypti GN=AAEL004243 PE=4 SV=1
CCG015615.1	1899	75	941	1.69706693	17.32266374	3.351545426	Up	0	0	AAEL012360-PA OS=Aedes aegypti GN=AAEL012360 PE=4 SV=1
CCG023684.1	2954	97	941	1.41098993	11.13599812	2.980451274	Up	3.62E-14	1.25E-13	Ubiquitin carboxyl-terminal hydrolase OS=Culex quinquefasciatus GN=CpipJ_CPIJ010867 PE=3 SV=1
CCG018660.1	2520	20	942	0.34102964	13.06773679	5.259968359	Up	3.69E-10	7.72E-10	AAEL004302-PA OS=Aedes aegypti GN=AAEL004302 PE=4 SV=1
CCG020469.1	981	217	942	9.50502794	33.56849819	1.820345221	Up	1.47E-13	4.37E-13	AAEL002644-PA OS=Aedes aegypti GN=AAEL002644 PE=4 SV=1
CCG023270.1	885	239	942	11.604256	37.2098268	1.681029645	Up	6.72E-13	1.70E-12	Proteasome subunit beta type OS=Aedes aegypti GN=AAEL006174 PE=3 SV=1
CCG002796.1	4108	321	942	3.35766427	8.016235813	1.255466966	Up	0	0	AAEL006087-PA OS=Aedes aegypti GN=AAEL006087 PE=4 SV=1
CCG002440.1	3324	175	943	2.26224535	9.917465402	2.132216053	Up	3.55E-13	9.58E-13	AAEL011147-PA OS=Aedes aegypti GN=AAEL011147 PE=4 SV=1
CCG018844.1	4668	256	943	2.35652358	7.062051199	1.583427164	Up	0	0	AAEL006790-PA OS=Aedes aegypti GN=AAEL006790 PE=4 SV=1
CCG019683.1	3987	314	943	3.38412257	8.268285678	1.288806416	Up	4.97E-14	1.68E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015044 PE=4 SV=1
CCG014192.1	2229	2447	943	47.1722479	14.78943697	-1.673371222	Down	1.46E-227	8.89E-227	AAEL005644-PA OS=Aedes aegypti GN=CLIFE12 PE=3 SV=1
CCG008409.1	3712	233	944	2.69718431	8.890251421	1.720770109	Up	0	0	AAEL003089-PA OS=Aedes aegypti GN=AAEL003089 PE=4 SV=1
CCG005734.1	1611	339	944	9.04204844	20.484552	1.17981479	Up	1.30E-13	3.98E-13	AAEL002384-PA OS=Aedes aegypti GN=AAEL002384 PE=4 SV=1

CCG024231.1	1566	165	945	4.52746246	21.09551185	2.220161509	Up	1.31E-14	4.67E-14	AAEL005476-PA OS=Aedes aegypti GN=AAEL005476 PE=4 SV=1
CCG016094.1	13783	194	945	0.60481234	2.396834619	1.986570881	Up	5.97E-13	1.53E-12	AAEL006715-PA OS=Aedes aegypti GN=AAEL006715 PE=4 SV=1
CCG025434.1	963	64	946	2.85572483	34.34115247	3.588009577	Up	0	0	AAEL008942-PA OS=Aedes aegypti GN=AAEL008942 PE=4 SV=1
CCG011602.1	2862	227	946	3.40815156	11.55504187	1.76146109	Up	0	0	AAEL005025-PA OS=Aedes aegypti GN=AAEL005025 PE=4 SV=1
CCG003440.1	2027	2846	946	60.3314577	16.31501225	-1.886710369	Down	0	0	AAEL014895-PA OS=Aedes aegypti GN=CYP6BB2 PE=3 SV=1
CCG014810.1	939	96	947	4.39307191	35.25611087	3.004571319	Up	0	0	AAEL006727-PA OS=Aedes aegypti GN=AAEL006727 PE=4 SV=1
CCG020186.1	870	3742	947	184.819249	38.05228518	-2.280060024	Down	0	0	AAEL010330-PA OS=Aedes aegypti GN=AAEL010330 PE=4 SV=1
CCG021603.1	3000	94	948	1.34638502	11.04681546	3.036467601	Up	0	0	AAEL004946-PA (Fragment) OS=Aedes aegypti GN=AAEL004946 PE=4 SV=1
CCG007523.1	1734	107	948	2.65153495	19.11213747	2.849589466	Up	0	0	AAEL000299-PA (Fragment) OS=Aedes aegypti GN=AAEL000299 PE=4 SV=1
CCG024234.1	411	219	948	22.8962819	80.63368949	1.816269393	Up	4.32E-13	1.14E-12	AAEL010634-PA OS=Aedes aegypti GN=AAEL010634 PE=4 SV=1
CCG011947.1	2334	227	948	4.17914728	14.19899159	1.764507965	Up	0	0	AAEL009424-PA OS=Aedes aegypti GN=AAEL009424 PE=4 SV=1
CCG012888.1	1971	263	948	5.73365814	16.81402658	1.552137463	Up	1.40E-14	4.98E-14	AAEL009502-PA OS=Aedes aegypti GN=AAEL009502 PE=4 SV=1
CCG016883.1	714	358	948	21.545049	46.41519101	1.107240675	Up	1.30E-13	3.97E-13	AAEL006699-PA OS=Aedes aegypti GN=AAEL006699 PE=4 SV=1
CCG004015.1	7005	8787	948	53.9007934	4.730970218	-3.510098526	Down	0	0	AAEL002204-PA OS=Aedes aegypti GN=AAEL002204 PE=4 SV=1
CCG017183.1	2334	167	949	3.07452686	14.21396943	2.208873188	Up	1.62E-13	4.77E-13	AAEL013597-PA OS=Aedes aegypti GN=AAEL013597 PE=4 SV=1
CCG002470.1	960	177	949	7.92254482	34.55771318	2.124971931	Up	0	0	AAEL008755-PA OS=Aedes aegypti GN=AAEL008755 PE=4 SV=1

CCG009367.1	2218	2735	949	52.9856737	14.95735106	-1.824747637	Down	5.03E-287	3.15E-286	AGAP004519-PA OS=Anopheles gambiae GN=AGAP004519 PE=4 SV=4
CCG009537.1	676	174	950	11.0602571	49.12775582	2.151153411	Up	1.78E-13	5.20E-13	AAEL006502-PA OS=Aedes aegypti GN=AAEL006502 PE=4 SV=1
CCG021889.1	1131	119	951	4.52113034	29.3946253	2.700796971	Up	0	0	AAEL000013-PA OS=Aedes aegypti GN=AAEL000013 PE=4 SV=1
CCG023593.1	2844	321	951	4.8499595	11.68963474	1.269185248	Up	0	0	AAEL003562-PA OS=Aedes aegypti GN=AAEL003562 PE=4 SV=1
CCG017038.1	1394	70	952	2.15773416	23.87394511	3.46784795	Up	0	0	AAEL006298-PA (Fragment) OS=Aedes aegypti GN=AAEL006298 PE=4 SV=1
CCG010423.1	714	181	952	10.8928879	46.6110357	2.09728508	Up	1.53E-13	4.55E-13	AAEL014098-PA OS=Aedes aegypti GN=AAEL014098 PE=4 SV=1
CCG027981.1	1870	318	952	7.30715273	17.7969409	1.284248012	Up	1.20E-13	3.69E-13	AAEL013982-PA OS=Aedes aegypti GN=AAEL013982 PE=4 SV=1
CCG001259.1	16478	1651	952	4.30531811	2.019679542	-1.091993438	Down	1.11E-80	5.64E-80	AAEL004242-PA OS=Aedes aegypti GN=AAEL004242 PE=4 SV=1
CCG010322.1	1224	166	953	5.82759473	27.21833151	2.223606176	Up	1.83E-13	5.31E-13	AAEL014376-PA (Fragment) OS=Aedes aegypti GN=AAEL014376 PE=4 SV=1
CCG007366.2	2328	306	953	5.64808367	14.31066914	1.341257765	Up	0	0	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1
CCG019394.1	2340	327	953	6.00474496	14.2372811	1.245498782	Up	6.36E-13	1.62E-12	AAEL007353-PA OS=Aedes aegypti GN=AAEL007353 PE=3 SV=1
CCG018386.1	1851	93	954	2.15893318	18.01739386	3.060999849	Up	4.04E-14	1.39E-13	AAEL006184-PA OS=Aedes aegypti GN=AAEL006184 PE=4 SV=1
CCG027746.1	717	158	954	9.46892339	46.51352307	2.296377912	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013057 PE=4 SV=1
CCG024679.1	909	218	954	10.3051729	36.68888453	1.831974335	Up	0	0	Proteasome subunit alpha type OS=Aedes aegypti GN=AAEL006061 PE=3 SV=1
CCG022489.1	949	220	954	9.96137156	35.14246158	1.818798946	Up	4.61E-13	1.20E-12	AAEL011665-PA OS=Aedes aegypti GN=AAEL011665 PE=2 SV=1

CCG024419.1	615	282	954	19.7031954	54.22796104	1.460607307	Up	5.45E-13	1.40E-12	AAEL003550-PA OS=Aedes aegypti GN=AAEL003550 PE=2 SV=1
CCG019132.3	5214	111	955	0.91477571	6.402983183	2.80725426	Up	0	0	AAEL002319-PA OS=Aedes aegypti GN=AAEL002319 PE=4 SV=1
CCG014529.1	1920	111	955	2.48418778	17.38810121	2.80725426	Up	0	0	Zinc finger protein 136 (Fragment) OS=Culex quinquefasciatus GN=CpipJ_CPIJ009500 PE=4 SV=1
CCG004451.1	1106	135	955	5.24494952	30.18549215	2.52485453	Up	1.90E-13	5.48E-13	AAEL002406-PA OS=Aedes aegypti GN=AAEL002406 PE=4 SV=1
CCG010556.1	1080	237	955	9.42946954	30.91217992	1.712926878	Up	2.32E-13	6.62E-13	AAEL008257-PA OS=Aedes aegypti GN=AAEL008257 PE=2 SV=1
CCG004445.1	2571	312	955	5.21453022	12.98527978	1.316267908	Up	9.28E-14	2.93E-13	AAEL001824-PA OS=Aedes aegypti GN=AAEL001824 PE=4 SV=1
CCG024792.1	1310	77	956	2.52570196	25.51153633	3.336393471	Up	0	0	AAEL013179-PA OS=Aedes aegypti GN=AAEL013179 PE=4 SV=1
CCG019892.1	2877	267	956	3.98780644	11.61630608	1.54248408	Up	5.21E-13	1.35E-12	AAEL006466-PA (Fragment) OS=Aedes aegypti GN=AAEL006466 PE=4 SV=1
CCG008417.1	1890	274	956	6.22947475	17.68259926	1.505147929	Up	0	0	AAEL003065-PA OS=Aedes aegypti GN=AAEL003065 PE=4 SV=1
CCG024406.1	1563	125	957	3.4364791	21.40439595	2.638904034	Up	2.35E-14	8.23E-14	AAEL002744-PA OS=Aedes aegypti GN=AAEL002744 PE=4 SV=1
CCG022363.1	3068	226	957	3.1653064	10.90452114	1.784509356	Up	2.04E-14	7.17E-14	AAEL006426-PA OS=Aedes aegypti GN=AAEL006426 PE=4 SV=1
CCG003610.1	2364	262	957	4.76229715	14.15189123	1.571265317	Up	0	0	Ubiquitin conjugating enzyme 7 interacting protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017240 PE=4 SV=1
CCG003990.1	1090	376	957	14.8225874	30.69272557	1.050099467	Up	3.71E-14	1.28E-13	Probable ER retained protein OS=Aedes aegypti PE=2 SV=1
CCG019744.1	1085	382	957	15.1285149	30.8341667	1.02725949	Up	6.17E-13	1.57E-12	AAEL008594-PA OS=Aedes aegypti GN=AAEL008594 PE=4 SV=1
CCG020699.1	1965	93	958	2.0336821	17.04327183	3.067036238	Up	4.04E-14	1.38E-13	AAEL009724-PA OS=Aedes aegypti GN=AAEL009724 PE=4 SV=1

CCG023101.1	1989	158	958	3.41338264	16.83762149	2.302414301	Up	0	0	AAEL001506-PA OS=Aedes aegypti GN=AAEL001506 PE=4 SV=1
CCG024514.1	3309	197	958	2.55818607	10.12089125	1.98414323	Up	0	0	AAEL004911-PA OS=Aedes aegypti GN=AAEL004911 PE=4 SV=1
CCG010140.1	1509	8	959	0.22780509	22.21669147	6.607700209	Up	2.19E-06	3.96E-06	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG014729.1	1854	98	959	2.27132362	18.08251749	2.992990365	Up	0	0	AAEL001124-PA OS=Aedes aegypti GN=AAEL001124 PE=4 SV=1
CCG027261.1	2791	349	959	5.3731413	12.01181921	1.160616983	Up	4.55E-14	1.54E-13	AAEL001317-PA OS=Aedes aegypti GN=AAEL001317 PE=3 SV=1
CCG009748.1	2328	2063	959	38.0784203	14.40076779	-1.402827897	Down	2.75E-148	1.56E-147	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011916 PE=4 SV=1
CCG009928.1	1371	176	960	5.51617308	24.47844325	2.149772181	Up	0	0	AAEL013071-PA OS=Aedes aegypti GN=AAEL013071 PE=4 SV=1
CCG001468.1	1545	1962	960	54.5673911	21.7216477	-1.328905527	Down	5.70E-130	3.17E-129	AAEL006514-PA OS=Aedes aegypti GN=AAEL006514 PE=4 SV=1
CCG011677.1	2907	127	961	1.87724675	11.55655452	2.622021138	Up	1.57E-13	4.67E-13	AAEL004364-PA OS=Aedes aegypti GN=AAEL004364 PE=4 SV=1
CCG017946.1	2371	256	961	4.63949897	14.16908645	1.610705825	Up	0	0	AAEL002258-PA OS=Aedes aegypti GN=AAEL002258 PE=4 SV=1
CCG006482.1	2592	261	961	4.32681355	12.96099691	1.582799828	Up	8.02E-14	2.58E-13	AAEL008807-PA OS=Aedes aegypti GN=AAEL008807 PE=4 SV=1
CCG008513.1	1823	129	962	3.04064496	18.44753826	2.600979032	Up	0	0	AAEL004232-PA OS=Aedes aegypti GN=AAEL004232 PE=4 SV=1
CCG002399.1	1230	254	962	8.87342487	27.34135143	1.623521601	Up	2.35E-13	6.67E-13	AAEL005822-PA OS=Aedes aegypti GN=AAEL005822 PE=3 SV=1
CCG020654.1	1135	297	962	11.2440627	29.62983459	1.397887167	Up	2.44E-13	6.86E-13	AAEL010083-PA OS=Aedes aegypti GN=IMD PE=4 SV=1
CCG028231.1	2283	6197	962	116.637514	14.73055727	-2.985147965	Down	0	0	AAEL012556-PA OS=Aedes aegypti GN=AAEL012556 PE=4 SV=1
CCG007505.1	7950	2327	963	12.5774305	4.23456862	-1.570550304	Down	1.50E-197	8.96E-197	-
CCG025222.1	3657	122	964	1.43349949	9.215143234	2.684465202	Up	8.04E-14	2.58E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_17442 PE=4 SV=1

CCG006643.1	1961	146	964	3.19917453	17.18499684	2.425377981	Up	9.33E-15	3.34E-14	AAEL007368-PA OS=Aedes aegypti GN=AAEL007368 PE=4 SV=1
CCG022266.1	1867	203	965	4.67212433	18.06895398	1.951362419	Up	9.28E-14	2.94E-13	AAEL003277-PA OS=Aedes aegypti GN=AAEL003277 PE=3 SV=1
CCG027653.1	1097	343	965	13.4353865	30.75181138	1.19463357	Up	0	0	AAEL013212-PA OS=Aedes aegypti GN=AAEL013212 PE=4 SV=1
CCG023155.1	5906	55	966	0.40015838	5.717862405	3.836832869	Up	0	0	AAEL008123-PA OS=Aedes aegypti GN=AAEL008123 PE=4 SV=1
CCG018364.1	499	78	966	6.71671202	67.6747402	3.332790364	Up	0	0	AAEL008719-PA OS=Aedes aegypti GN=AAEL008719 PE=4 SV=1
CCG025512.1	1185	148	966	5.36668415	28.49763322	2.408739217	Up	0	0	AAEL009894-PA OS=Aedes aegypti GN=AAEL009894 PE=4 SV=1
CCG022422.4	1851	167	966	3.87679399	18.24402775	2.23448829	Up	1.62E-13	4.77E-13	AAEL004874-PA OS=Aedes aegypti GN=AAEL004874 PE=4 SV=1
CCG006880.1	10375	1838	966	7.61237323	3.254910396	-1.225728469	Down	1.46E-107	7.80E-107	AAEL007364-PA OS=Aedes aegypti GN=AAEL007364 PE=4 SV=1
CCG006827.1	1856	172	967	3.98210903	18.21371425	2.193420529	Up	1.35E-13	4.09E-13	Glyz1n OS=Culex quinquefasciatus
CCG001015.1	9023	191	967	0.90958875	3.746498242	2.042256455	Up	6.68E-14	2.19E-13	CN_CpipJ_CPIJ015048 PE=4 SV=1
CCG004839.1	2175	240	967	4.74148796	15.54236949	1.712794688	Up	0	0	AAEL010101-PA (Fragment) OS=Aedes aegypti GN=AAEL010101 PE=4 SV=1
CCG027614.1	475	298	967	26.9578546	71.16769187	1.400516763	Up	3.92E-13	1.05E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002917 PE=4 SV=1
CCG009175.1	4005	1842	967	19.7628592	8.440612644	-1.227372063	Down	5.11E-108	2.74E-107	AAEL005319-PA OS=Aedes aegypti GN=AAEL005319 PE=4 SV=1
CCG012095.1	957	286	968	12.8415299	35.36009605	1.461305104	Up	0	0	AAEL010099-PA OS=Aedes aegypti GN=AAEL010099 PE=4 SV=1
CCG025958.1	962	378	968	16.8841577	35.17631176	1.058934017	Up	1.08E-12	2.65E-12	AAEL012276-PA OS=Aedes aegypti GN=AAEL012276 PE=2 SV=1
CCG027654.1	1054	305	969	12.434316	32.1390609	1.370000627	Up	5.30E-13	1.37E-12	AAEL013207-PA OS=Aedes aegypti GN=AAEL013207 PE=4 SV=1
CCG002902.1	1862	191	971	4.40774399	18.23012178	2.048211861	Up	6.68E-14	2.20E-13	AAEL012137-PA OS=Aedes aegypti GN=AAEL012137 PE=4 SV=1

CCG014286.1	14727	1636	971	4.77344237	2.30491524	-1.050316344	Down	3.92E-75	1.97E-74	AAEL014378-PA OS=Aedes aegypti GN=AAEL014378 PE=4 SV=1
CCG004665.1	3522	26	972	0.31720985	9.64776974	4.926685989	Up	4.60E-12	1.05E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_17442 PE=4 SV=1
CCG022393.1	693	38	972	2.35620478	49.0323882	4.379198194	Up	4.26E-14	1.45E-13	AAEL004492-PA OS=Aedes aegypti GN=AAEL004492 PE=4 SV=1
CCG006785.1	1722	85	972	2.121038	19.73254647	3.217734771	Up	0	0	AAEL008878-PA OS=Aedes aegypti GN=AAEL008878 PE=4 SV=1
CCG014790.1	2076	320	972	6.62346584	16.36774808	1.305197613	Up	0	0	AAEL002651-PA OS=Aedes aegypti GN=AAEL002651 PE=4 SV=1
CCG006279.1	3554	358	972	4.32840883	9.560901807	1.14330993	Up	1.30E-13	3.97E-13	AAEL000965-PA OS=Aedes aegypti GN=AAEL000965 PE=4 SV=1
CCG000345.1	3620	1643	972	19.5025619	9.386587023	-1.054991058	Down	5.48E-76	2.76E-75	AAEL003698-PA OS=Aedes aegypti GN=AAEL003698 PE=4 SV=1
CCG022912.1	1667	1680	972	43.3048315	20.3835903	-1.08711981	Down	1.76E-81	8.96E-81	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012060 PE=4 SV=1
CCG025281.3	2760	2219	972	34.5470439	12.31139312	-1.488568245	Down	3.64E-174	2.12E-173	AAEL002043-PA OS=Aedes aegypti GN=CYP305A5 PE=3 SV=1
CCG027366.1	1104	332	973	12.9220579	30.81014792	1.253569767	Up	0	0	AAEL005259-PA OS=Aedes aegypti GN=AAEL005259 PE=4 SV=1
CCG007122.1	1407	206	974	6.29123335	24.19997269	1.943590639	Up	1.10E-13	3.43E-13	AAEL007778-PA OS=Aedes aegypti GN=AAEL007778 PE=4 SV=1
CCG013247.1	849	20	975	1.01224345	40.14643092	5.309643518	Up	3.69E-10	7.72E-10	AAEL008878-PA OS=Aedes aegypti GN=AAEL008878 PE=4 SV=1
CCG008843.1	2493	100	975	1.72361551	13.67200957	2.987715423	Up	0	0	AAEL007512-PA OS=Aedes aegypti GN=AAEL007512 PE=4 SV=1
CCG000140.1	2835	129	975	1.9552366	12.02268778	2.620344357	Up	0	0	AAEL008333-PA OS=Aedes aegypti GN=AAEL008333 PE=4 SV=1
CCG027008.1	2292	223	975	4.1807377	14.8709947	1.830671713	Up	0	0	AAEL007638-PA OS=Aedes aegypti GN=AAEL007638 PE=4 SV=1
CCG014509.1	2337	287	975	5.27698495	14.58464692	1.466664686	Up	2.87E-13	7.98E-13	-

CCG001108.1	8482	331	975	1.67684298	4.018429598	1.260884206	Up	0	0	Numo-associated kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ004075 PE=4 SV=1
CCG001347.1	1519	393	975	11.1172519	22.43865691	1.01318611	Up	1.08E-13	3.37E-13	AAEL014141-PA OS=Aedes aegypti GN=SRPN5 PE=3 SV=1
CCG007110.2	2952	3103	975	45.1677122	11.54617881	-1.967876369	Down	0	0	AAEL005187-PA OS=Aedes aegypti GN=AAEL005187 PE=4 SV=1
CCG013858.1	1158	30	976	1.11320556	29.46397075	4.726159946	Up	3.71E-13	9.95E-13	AAEL006803-PA (Fragment) OS=Aedes aegypti GN=AAEL006803 PE=4 SV=1
CCG028535.1	4775	158	976	1.42182577	7.145398561	2.329269793	Up	0	0	AAEL004571-PA OS=Aedes aegypti GN=AAEL004571 PE=4 SV=1
CCG015427.1	2750	162	976	2.531308	12.40701023	2.293200538	Up	0	0	AAEL007729-PA OS=Aedes aegypti GN=AAEL007729 PE=4 SV=1
CCG001707.1	9325	1766	976	8.13775349	3.658903821	-1.153219086	Down	5.61E-94	2.94E-93	AAEL010047-PA OS=Aedes aegypti GN=AAEL010047 PE=3 SV=1
CCG027015.1	1992	2026	976	43.7031538	17.12815167	-1.351367917	Down	1.25E-137	7.03E-137	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001433 PE=4 SV=1
CCG006374.1	1323	90	977	2.9231112	25.81574936	3.142674859	Up	0	0	histone acetyltransferase Tipou OS=Culex quinquefasciatus GN=CpipJ_CPIJ001206 PE=4 SV=1
CCG020201.1	1194	319	977	11.4801887	28.60488811	1.317115342	Up	0	0	AAEL008084-PA OS=Aedes aegypti GN=AAEL008084 PE=4 SV=1
CCG013993.1	1965	31	978	0.67789403	17.39908126	4.681807548	Up	2.40E-13	6.77E-13	AAEL007692-PA OS=Aedes aegypti GN=AAEL007692 PE=4 SV=1
CCG022281.1	2922	191	978	2.80876773	11.7006142	2.058575031	Up	6.68E-14	2.19E-13	-
CCG010759.1	1089	260	978	10.2590735	31.39503644	1.613636046	Up	3.22E-13	8.81E-13	AAEL005145-PA OS=Aedes aegypti GN=AAEL005145 PE=4 SV=1
CCG013323.1	3058	1880	978	26.4169722	11.18024679	-1.240513088	Down	4.72E-112	2.55E-111	AAEL007658-PA (Fragment) OS=Aedes aegypti GN=AAEL007658 PE=4 SV=1
CCG007310.1	6326	3669	978	24.9219027	5.404551799	-2.20516733	Down	0	0	AAEL014026-PA OS=Aedes aegypti GN=AAEL014026 PE=4 SV=1

CCG013161.1	1911	28	979	0.62959318	17.90902824	4.830123331	Up	1.17E-12	2.85E-12	AAEL004049-PA OS=Aedes aegypti GN=AAEL004049 PE=4 SV=1
CCG027509.1	1737	49	979	1.21215717	19.70302416	4.022768409	Up	0	0	AAEL013842-PA OS=Aedes aegypti GN=AAEL013842 PE=4 SV=1
CCG017134.1	1470	78	979	2.28002673	23.28173671	3.352076035	Up	0	0	AAEL009990-PA OS=Aedes aegypti GN=AAEL009990 PE=4 SV=1
CCG004228.1	2909	113	979	1.66915779	11.76492023	2.817299291	Up	1.42E-13	4.26E-13	Ribonuclease P/MKP protein subunit POP1 OS=Culex quinquefasciatus
CCG015047.1	2198	129	979	2.5218816	15.57058824	2.626250998	Up	0	0	AAEL011072-PA OS=Aedes aegypti GN=AAEL011072 PE=4 SV=1
CCG017857.1	1152	263	979	9.80993073	29.70846611	1.598559264	Up	1.40E-14	4.97E-14	Blood meal-induced protein OS=Aedes aegypti GN=MRRG PE=2 SV=2
CCG000338.1	11411	355	980	1.33680272	3.002288251	1.167275928	Up	2.28E-13	6.52E-13	AAEL003652-PA OS=Aedes aegypti GN=AAEL003652 PE=3 SV=1
CCG012140.1	2541	1618	980	27.3612871	13.48253099	-1.02104475	Down	5.10E-71	2.53E-70	AAEL011967-PA OS=Aedes aegypti GN=AAEL011967 PE=4 SV=1
CCG002107.2	29945	1770	980	2.53987077	1.144067832	-1.150582502	Down	7.59E-94	3.97E-93	-
CCG028516.1	4796	4405	980	39.4665723	7.143267564	-2.465975161	Down	0	0	AAEL002815-PA OS=Aedes aegypti GN=AAEL002815 PE=4 SV=1
CCG026458.1	1692	121	982	3.07289473	20.28902352	2.723029181	Up	0	0	AAEL008000-PA OS=Aedes aegypti GN=AAEL008000 PE=4 SV=1
CCG014478.2	1920	149	982	3.33463045	17.87970197	2.422723898	Up	2.33E-13	6.63E-13	Heat shock 70 Ab OS=Aedes aegypti GN=Hsp70Ab PE=3 SV=1
CCG012849.1	4054	283	982	2.99961394	8.467939761	1.497234175	Up	0	0	AAEL006062-PA (Fragment) OS=Aedes aegypti GN=AAEL006062 PE=4 SV=1
CCG017196.2	1637	138	984	3.62237225	21.01340522	2.536303252	Up	1.49E-13	4.45E-13	AAEL012787-PA OS=Aedes aegypti GN=AAEL012787 PE=4 SV=1
CCG012324.1	666	158	984	10.1940211	51.65006658	2.341046961	Up	0	0	AAEL015143-PA OS=Aedes aegypti GN=AAEL015143 PE=2 SV=1
CCG011480.1	2104	71	985	1.45002431	16.3659233	3.496545999	Up	0	0	AAEL000904-PA (Fragment) OS=Aedes aegypti GN=AAEL000904 PE=4 SV=1

CCG022779.1	1017	83	985	3.50687116	33.85831133	3.271253687	Up	1.32E-13	4.01E-13	AAEL011023-PA OS=Aedes aegypti GN=AAEL011023 PE=4 SV=1
CCG028171.1	771	291	985	16.2181489	44.6613523	1.461417775	Up	6.00E-14	1.99E-13	AAEL0006904-PA OS=Aedes aegypti GN=AAEL006904 PE=4 SV=1
CCG002367.1	678	5788	985	366.827174	50.78746699	-2.852556088	Down	0	0	AAEL0009076-PA OS=Aedes aegypti GN=AAEL009078 PE=4 SV=1
CCG026225.1	375	10785	985	1235.80957	91.82374032	-3.750445438	Down	0	0	AAEL007306-PA OS=Aedes aegypti GN=AAEL007306 PE=4 SV=1
CCG019741.1	1245	121	987	4.17617501	27.71391098	2.730356241	Up	0	0	AAEL008599-PA (Fragment) OS=Aedes aegypti GN=AAEL008599 PE=3 SV=1
CCG018507.1	831	166	987	8.58360523	41.52084136	2.274180047	Up	1.83E-13	5.32E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009943 PE=4 SV=1
CCG024353.1	2479	190	987	3.29336409	13.91844259	2.07936387	Up	2.69E-13	7.52E-13	Zinc finger protein z6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ016622 PE=4 SV=1
CCG016675.1	2808	226	987	3.45839032	12.28768489	1.829040516	Up	2.04E-14	7.16E-14	AAEL000440-PA OS=Aedes aegypti GN=AAEL000440 PE=4 SV=1
CCG003495.1	630	257	988	17.5289235	54.82345627	1.645055886	Up	4.14E-13	1.10E-12	AAEL0009317-PA OS=Aedes aegypti GN=AAEL009317 PE=2 SV=1
CCG005864.1	2151	102	989	2.03761643	16.07333135	2.979714573	Up	0	0	AAEL013993-PA OS=Aedes aegypti GN=AAEL013993 PE=4 SV=1
CCG018268.1	1080	130	989	5.17228287	32.01271827	2.629772102	Up	0	0	AAEL009739-PA OS=Aedes aegypti GN=AAEL009739 PE=4 SV=1
CCG016004.1	1149	225	989	8.41443889	30.09028349	1.838358723	Up	0	0	AAEL001389-PA OS=Aedes aegypti GN=AAEL001389 PE=4 SV=1
CCG014541.1	2298	47	990	0.8788414	15.06035422	4.099009067	Up	0	0	AAEL009799-PA OS=Aedes aegypti GN=AAEL009799 PE=4 SV=1
CCG000783.1	1752	301	990	7.38235737	19.75382078	1.419978242	Up	0	0	AAEL000594-PA OS=Aedes aegypti GN=AAEL000594 PE=4 SV=1
CCG016093.1	1136	319	990	12.0663251	30.46539965	1.336185305	Up	0	0	Splicing factor Uzar 56 kDa subunit OS=Acromyrmex echinator GN=G5I_01763 PE=4 SV=1

CCG008524.1	2004	17	991	0.36451372	17.28725164	5.56759161	Up	3.39E-09	6.90E-09	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002013 PE=4 SV=1
CCG018607.1	1095	30	991	1.177253	31.63803861	4.748163855	Up	3.71E-13	9.94E-13	AAEL005000-PA OS=Aedes aegypti GN=AAEL005000 PE=4 SV=1
CCG021839.2	1779	402	991	9.70985571	19.47366626	1.00400276	Up	0	0	AAEL006139-PA OS=Aedes aegypti GN=AAEL006139 PE=3 SV=1
CCG019980.1	1638	74	992	1.94124564	21.17131292	3.447056149	Up	0	0	AAEL005585-PA OS=Aedes aegypti GN=AAEL005585 PE=4 SV=1
CCG022973.1	1592	288	992	7.77341933	21.78304683	1.486584513	Up	0	0	AAEL002521-PA OS=Aedes aegypti GN=AAEL002521 PE=4 SV=1
CCG017482.1	1365	57	993	1.79434057	25.43118596	3.825073097	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_13527 PE=4 SV=1
CCG010713.1	2019	64	993	1.36209164	17.19344667	3.657963111	Up	0	0	AAEL003792-PA OS=Aedes aegypti GN=AAEL003792 PE=4 SV=1
CCG001854.1	1353	124	993	3.93809837	25.65673972	2.703766801	Up	1.61E-13	4.77E-13	AAEL013531-PA OS=Aedes aegypti GN=AAEL013531 PE=3 SV=1
CCG005130.1	1356	260	993	8.23903466	25.59997702	1.635595298	Up	3.22E-13	8.81E-13	AAEL006321-PA OS=Aedes aegypti GN=AAEL006321 PE=4 SV=1
CCG003232.1	1278	293	993	9.85143368	27.16241693	1.463206257	Up	4.19E-13	1.11E-12	AAEL004183-PA OS=Aedes aegypti GN=AAEL004183 PE=4 SV=1
CCG015619.1	2547	174	994	2.93550602	13.64292388	2.21647175	Up	1.78E-13	5.20E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015878 PE=4 SV=1
CCG011509.1	1080	230	994	9.150962	32.17456214	1.813925194	Up	3.85E-13	1.03E-12	AAEL007066-PA OS=Aedes aegypti GN=AAEL007066 PE=4 SV=1
CCG011323.1	1644	74	995	1.9341608	21.15783783	3.451412554	Up	0	0	Venus kinase receptor (Fragment) OS=Aedes aegypti GN=VKR PE=2 SV=1
CCG003390.1	1725	241	995	6.00330785	20.16433936	1.747976583	Up	2.37E-13	6.72E-13	AAEL010107-PA OS=Aedes aegypti GN=AAEL010107 PE=4 SV=1
CCG027397.1	662	263	995	17.0710577	52.54302929	1.62194693	Up	1.40E-14	4.97E-14	AAEL000847-PA OS=Aedes aegypti GN=AAEL000847 PE=4 SV=1

CCG007955.1	2009	2550	995	54.5409772	17.31383046	-1.655415612	Down	2.04E-233	1.24E-232	AAEL005738-PA OS=Aedes aegypti GN=AAEL005738 PE=4 SV=1
CCG018728.1	2769	3153	995	48.9287011	12.56174987	-1.961643535	Down	0	0	AAEL000066-PA OS=Culex quinquefasciatus GN=CripJ_CPIJ001691 PE=4 SV=1
CCG013414.2	4551	353	996	3.33296337	7.650723723	1.198790763	Up	0	0	AAEL010888-PA OS=Aedes aegypti GN=AAEL010888 PE=4 SV=1
CCG005334.2	3141	394	996	5.3900272	11.08514603	1.040263316	Up	5.49E-13	1.41E-12	AAEL000070-PA OS=Aedes aegypti GN=AAEL000070 PE=4 SV=1
CCG027412.1	1833	3003	996	70.3972248	18.99533206	-1.889873624	Down	0	0	Synaptotagmin I isoform A OS=Culex pipiens GN=sytl PE=2 SV=1
CCG001008.2	1776	111	997	2.68560841	19.62466326	2.869347032	Up	0	0	AAEL002272-PA OS=Aedes aegypti GN=AAEL002272 PE=4 SV=1
CCG026826.1	2205	133	998	2.59182526	15.82238559	2.609926774	Up	9.90E-14	3.11E-13	AAEL009912-PA OS=Aedes aegypti GN=AAEL009912 PE=4 SV=1
CCG018831.1	2642	378	998	6.14782728	13.20528396	1.102966785	Up	1.08E-12	2.65E-12	AAEL000925-PA OS=Aedes aegypti GN=AAEL000925 PE=4 SV=1
CCG021607.1	1536	188	999	5.25931648	22.73653548	2.11206522	Up	1.80E-13	5.25E-13	U4/U6 small nuclear ribonucleoprotein Prp31 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002179 PE=4 SV=1
CCG015587.1	1749	198	999	4.86449826	19.96759205	2.037297452	Up	0	0	AAEL013211-PA OS=Aedes aegypti GN=AAEL013211 PE=4 SV=1
CCG004830.1	783	339	999	18.6037548	44.60193933	1.261512608	Up	1.30E-13	3.98E-13	AAEL010112-PA OS=Aedes aegypti GN=AAEL010112 PE=4 SV=1
CCG008788.2	938	3308	999	151.53932	37.23168283	-2.025089448	Down	0	0	AGAP005507-PA OS=Anopheles gambiae GN=AGAP005507 PE=4 SV=1
CCG001576.1	1317	208	1000	6.78641215	26.54386999	1.96765777	Up	3.93E-13	1.05E-12	AAEL001371-PA OS=Aedes aegypti GN=AAEL001371 PE=4 SV=1
CCG005089.1	2883	237	1000	3.53237152	12.12565965	1.77935424	Up	2.32E-13	6.61E-13	Lon protease homolog OS=Aedes aegypti GN=AAEL006474 PE=3 SV=1
CCG012247.1	1437	3207	1000	95.8969651	24.32726289	-1.97891115	Down	0	0	AAEL003128-PA OS=Aedes aegypti GN=AAEL003128 PE=4 SV=1

CCG005053.1	1287	141	1001	4.70763992	27.18977082	2.52998811	Up	0	0	AAEL012426-PA OS=Aedes aegypti GN=AAEL012426 PE=4 SV=1
CCG015188.1	1221	213	1001	7.49594879	28.65948816	1.934829842	Up	3.57E-13	9.60E-13	AAEL006113-PA OS=Aedes aegypti GN=AAEL006113 PE=4 SV=1
CCG013876.1	2979	10780	1001	155.493031	11.74663815	-3.726530095	Down	0	0	Ubiquitin carboxyl-terminal hydrolase OS=Aedes aegypti GN=AAEL010854 PE=3 SV=1
CCG017229.1	1440	81	1002	2.41704757	24.32513425	3.331129994	Up	4.46E-14	1.51E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005826 PE=4 SV=1
CCG013192.1	1359	94	1002	2.97215236	25.77497669	3.116391145	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000820 PE=4 SV=1
CCG020324.1	1812	169	1002	4.00766289	19.33123252	2.270100561	Up	0	0	AAEL011267-PA OS=Aedes aegypti GN=AAEL011267 PE=4 SV=1
CCG023484.1	1539	262	1002	7.31518549	22.76035954	1.637556995	Up	0	0	AAEL011065-PA OS=Aedes aegypti GN=AAEL011065 PE=4 SV=1
CCG005760.1	1305	210	1003	6.91466994	26.8683154	1.958173577	Up	0	0	AAEL004443-PA OS=Aedes aegypti GN=AAEL004443 PE=4 SV=1
CCG004112.1	2007	48	1004	1.02767676	17.48784747	4.088894257	Up	5.35E-14	1.79E-13	AAEL001357-PA OS=Aedes aegypti GN=AAEL001357 PE=4 SV=1
CCG002105.1	483	191	1004	16.9921725	72.66689416	2.09642793	Up	6.68E-14	2.19E-13	AAEL003165-PA OS=Aedes aegypti GN=AAEL003165 PE=4 SV=1
CCG027668.1	17138	4156	1004	10.4202484	2.047970001	-2.347123181	Down	0	0	AAEL007619-PA OS=Aedes aegypti GN=TOLL5A PE=4 SV=1
CCG017327.1	1002	80	1005	3.43071733	35.06294227	3.353364895	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011214 PE=4 SV=1
CCG022306.1	2343	7465	1005	136.905279	14.99490745	-3.190633555	Down	0	0	AAEL011094-PA OS=Aedes aegypti GN=AAEL011094 PE=4 SV=1
CCG008383.1	2181	133	1006	2.62034604	16.12472555	2.621445358	Up	9.90E-14	3.11E-13	AAEL010176-PA OS=Aedes aegypti GN=AAEL010176 PE=4 SV=1
CCG025103.1	1320	193	1006	6.28269605	26.64244427	2.084270756	Up	6.08E-14	2.02E-13	AAEL002107-PA OS=Aedes aegypti GN=AAEL002107 PE=4 SV=1

CCG005357.1	2307	278	1006	5.17797409	15.24405134	1.557786721	Up	2.11E-14	7.39E-14	AAEL010800-PA OS=Aedes aegypti GN=AAEL010800 PE=4 SV=1
CCG018481.1	579	249	1007	18.4792123	60.79962817	1.71815924	Up	0	0	AAEL010504-PA OS=Aedes aegypti GN=AAEL010504 PE=4 SV=1
CCG016127.1	2576	295	1007	4.92083529	13.66575493	1.473590028	Up	2.32E-13	6.61E-13	Dicer 2 isoform A OS=Aedes albopictus GN=Dcr-2 PE=2 SV=1
CCG002362.1	1674	138	1008	3.54230787	21.05014515	2.571068671	Up	1.49E-13	4.45E-13	AAEL002249-PA OS=Aedes aegypti GN=AAEL002249 PE=4 SV=1
CCG002790.1	732	4229	1008	248.25001	48.13926637	-2.366507718	Down	0	0	AAEL013744-PA OS=Aedes aegypti GN=AAEL013744 PE=2 SV=1
CCG027709.1	830	2190	1009	113.377974	42.4974714	-1.415691492	Down	1.79E-159	1.03E-158	AAEL005772-PA OS=Aedes aegypti GN=AAEL005772 PE=2 SV=1
CCG011025.2	2484	90	1010	1.55687444	14.21411415	3.190599685	Up	0	0	AAEL014265-PA OS=Aedes aegypti GN=AAEL014265 PE=4 SV=1
CCG025770.1	1032	136	1010	5.6626782	34.21304219	2.59498994	Up	1.70E-13	4.98E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_01799 PE=4 SV=1
CCG014127.1	2403	162	1010	2.89683604	14.69324159	2.342602779	Up	0	0	AAEL006032-PA OS=Aedes aegypti GN=AAEL006032 PE=4 SV=1
CCG007654.1	1959	361	1010	7.91836355	18.02340967	1.186597755	Up	1.35E-13	4.08E-13	AAEL006041-PA (Fragment) OS=Aedes aegypti GN=AAEL006041 PE=4 SV=1
CCG003060.1	1738	2209	1010	54.6145821	20.31522413	-1.426724922	Down	9.86E-163	5.69E-162	AAEL006910-PA OS=Aedes aegypti GN=AAEL006910 PE=4 SV=1
CCG019622.4	2202	411	1011	8.02023657	16.05032598	1.000885902	Up	0	0	Coronin OS=Aedes aegypti GN=AAEL001360 PE=3 SV=1
CCG002031.1	2081	139	1012	2.87015527	17.00037294	2.566365706	Up	1.46E-13	4.37E-13	AAEL010847-PA OS=Aedes aegypti GN=AAEL010847 PE=4 SV=1
CCG006138.1	1682	3610	1013	92.2239845	21.05394433	-2.131051459	Down	0	0	AAEL004951-PA OS=Aedes aegypti GN=AAEL004951 PE=4 SV=1
CCG023679.1	2038	262	1014	5.52407776	17.39337225	1.654732139	Up	0	0	AAEL006409-PA OS=Aedes aegypti GN=AAEL006409 PE=4 SV=1
CCG024808.1	2955	61	1015	0.88702329	12.00766529	3.758839878	Up	1.75E-14	6.18E-14	AAEL011329-PA (Fragment) OS=Aedes aegypti GN=AAEL011329 PE=4 SV=1

CCG024107.1	1023	131	1015	5.50247823	34.68489826	2.656154214	Up	1.26E-13	3.87E-13	AAEL008088-PA OS=Aedes aegypti GN=AAEL008088 PE=4 SV=1
CCG010509.1	759	395	1015	22.3623784	46.7492107	1.063868373	Up	0	0	AAEL001496-PA OS=Aedes aegypti GN=AAEL001496 PE=4 SV=1
CCG012443.1	979	2202	1015	96.6489843	36.2437701	-1.415021538	Down	3.19E-160	1.83E-159	AAEL011992-PA OS=Aedes aegypti GN=AAEL011992 PE=4 SV=1
CCG023897.1	743	49	1016	2.83380484	47.80297335	4.076288046	Up	0	0	-
CCG006554.2	3270	79	1016	1.03810674	10.86165419	3.387217142	Up	0	0	AAEL010010-PA OS=Aedes aegypti GN=AAEL010010 PE=4 SV=1
CCG016322.1	2124	157	1016	3.17619978	16.72203823	2.396377142	Up	0	0	AAEL007058-PA OS=Aedes aegypti GN=AAEL007058 PE=4 SV=1
CCG016654.1	1332	262	1017	8.45200486	26.69111672	1.658994166	Up	0	0	AAEL012440-PA OS=Aedes aegypti GN=AAEL012440 PE=4 SV=1
CCG014280.1	1701	42	1018	1.0609811	20.9215319	4.301517627	Up	7.66E-14	2.47E-13	AAEL015080-PA OS=Aedes aegypti GN=AAEL015080 PE=4 SV=1
CCG016437.1	1479	1674	1018	48.6351154	24.06188354	-1.015248763	Down	9.68E-73	4.84E-72	AAEL003697-PA OS=Aedes aegypti GN=SRPN17 PE=3 SV=2
CCG025560.1	4866	1998	1018	17.6435532	7.313507142	-1.270505818	Down	1.83E-123	1.01E-122	AAEL009266-PA OS=Aedes aegypti GN=AAEL009266 PE=4 SV=1
CCG025968.1	1119	235	1019	9.02402827	31.83421272	1.818734593	Up	4.34E-13	1.14E-12	AAEL002166-PA OS=Aedes aegypti GN=AAEL002166 PE=4 SV=1
CCG021664.1	930	289	1019	13.3529605	38.30374627	1.520325857	Up	0	0	Caspase OS=Culex quinquefasciatus GN=CpipJ_CPIJ000671 PE=3 SV=1
CCG003794.1	633	64	1020	4.34449134	56.33087252	3.696666641	Up	0	0	AAEL001955-PA OS=Aedes aegypti GN=AAEL001955 PE=4 SV=1
CCG003408.1	1356	1829	1020	57.95844	26.2960489	-1.140172719	Down	1.59E-95	8.35E-95	Carboxypeptidase A OS=Aedes aegypti GN=CPA-II PE=2 SV=1
CCG017967.1	1629	216	1021	5.69764437	21.91062037	1.943192853	Up	1.38E-13	4.17E-13	AAEL001173-PA OS=Aedes aegypti GN=AAEL001173 PE=4 SV=1
CCG018248.1	1353	406	1021	12.8940963	26.3801926	1.032744438	Up	1.21E-13	3.73E-13	Caspase OS=Aedes aegypti GN=CASPL2 PE=3 SV=1
CCG010996.1	1200	3729	1021	133.52845	29.74366715	-2.166492727	Down	0	0	AAEL002781-PA OS=Aedes aegypti GN=AAEL002781 PE=3 SV=1

CCG000945.1	645	0	1022	0.001	55.39125405	15.75737058	Up	4.44E-16	1.66E-15	AAEL008182-PA OS=Aedes aegypti GN=AAEL008182 PE=4 SV=1
CCG012088.1	1654	52	1022	1.35092273	21.60057972	3.999052967	Up	0	0	AAEL003763-PA (Fragment) OS=Aedes aegypti GN=CYP329B1 PE=3 SV=1
CCG008444.1	1785	73	1022	1.75730567	20.01532709	3.509668126	Up	0	0	Flizzled-5 OS=Culex quinquefasciatus GN=CpipJ_CPIJ018158 PE=4 SV=1
CCG006284.1	2364	73	1022	1.32689959	15.11309597	3.509668126	Up	0	0	AAEL000404-PA OS=Aedes aegypti GN=AAEL000404 PE=4 SV=1
CCG017986.1	2468	94	1022	1.63661064	14.47623941	3.144903833	Up	0	0	AAEL005719-PA OS=Aedes aegypti GN=AAEL005719 PE=4 SV=1
CCG007908.3	3058	187	1022	2.62764564	11.68324358	2.152598225	Up	3.31E-13	9.03E-13	AAEL005295-PA OS=Aedes aegypti GN=AAEL003293 PE=4 SV=1
CCG027370.1	2736	227	1022	3.56510591	13.0582452	1.872944197	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010636 PE=4 SV=1
CCG011192.2	4489	331	1022	3.1684077	7.958868091	1.328805278	Up	0	0	AAEL004790-PA OS=Aedes aegypti GN=AAEL004790 PE=4 SV=1
CCG007417.1	772	343	1022	19.0914754	46.27896226	1.277427919	Up	0	0	Nucleoside diphosphate kinase OS=Aedes aegypti GN=AAEL004107 PE=3 SV=1
CCG021142.1	1676	368	1022	9.43488206	21.31703989	1.175930729	Up	3.69E-13	9.91E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_17136 PE=4 SV=1
CCG010144.1	441	15662	1022	1526.05892	81.01441919	-4.235488148	Down	0	0	AAEL011184-PA OS=Aedes aegypti GN=AAEL011184 PE=3 SV=1
CCG001266.1	1595	224	1023	6.03462104	22.42151545	1.893548712	Up	2.04E-13	5.88E-13	AAEL003354-PA OS=Aedes aegypti GN=AAEL003354 PE=4 SV=1
CCG017156.1	2316	338	1023	6.27105799	15.441415	1.300024197	Up	1.44E-13	4.32E-13	Mediator complex OS=Culex quinquefasciatus GN=CpipJ_CPIJ001509 PE=4 SV=1
CCG010258.1	930	64	1024	2.95705701	38.49169399	3.702313204	Up	0	0	Eye pigment transporter OS=Aedes aegypti GN=white PE=2 SV=1

CCG007819.1	1095	211	1024	8.28001279	32.69157572	1.981214015	Up	1.30E-13	3.98E-13	AAEL010077-PA OS=Aedes aegypti GN=AAEL010077 PE=4 SV=1
CCG018361.1	2049	220	1024	4.61363671	17.47060782	1.92095349	Up	4.61E-13	1.20E-12	AAEL0008721-PA OS=Aedes aegypti GN=AAEL008721 PE=4 SV=1
CCG008448.1	4653	386	1024	3.56465024	7.693375331	1.109856167	Up	2.52E-13	7.09E-13	AAEL0008705-PA OS=Aedes aegypti GN=AAEL008705 PE=4 SV=1
CCG004525.1	746	6	1025	0.34560108	48.03248484	7.118758897	Up	8.38E-06	1.47E-05	AAEL012504-PA OS=Aedes aegypti GN=AAEL012504 PE=4 SV=1
CCG012405.1	1569	155	1025	4.24493873	22.83762504	2.427596993	Up	3.42E-13	9.25E-13	AAEL007328-PA OS=Aedes aegypti GN=AAEL007328 PE=4 SV=1
CCG020099.1	1305	163	1026	5.36710095	27.48443829	2.356400065	Up	1.80E-13	5.24E-13	AAEL012318-PA OS=Aedes aegypti GN=AAEL012318 PE=3 SV=1
CCG004002.1	1673	38	1027	0.97600114	21.45974312	4.458606157	Up	4.26E-14	1.45E-13	AGAP001938-PA OS=Anopheles gambiae GN=AgaP_AGAP001938 PE=4 SV=4
CCG024750.2	1836	198	1027	4.63399099	19.55454806	2.07717705	Up	0	0	AAEL011442-PA OS=Aedes aegypti GN=AAEL011442 PE=4 SV=1
CCG005616.1	1447	2405	1029	71.4182528	24.85975591	-1.522480708	Down	5.40E-195	3.21E-194	AAEL012311-PA OS=Aedes aegypti GN=AAEL012311 PE=3 SV=1
CCG027579.1	762	117	1030	6.59771516	47.25331375	2.840377106	Up	9.44E-14	2.97E-13	AAEL009579-PA OS=Aedes aegypti GN=AAEL009579 PE=4 SV=1
CCG005187.1	2085	119	1030	2.45246927	17.26955639	2.815924063	Up	0	0	AAEL003141-PA OS=Aedes aegypti GN=AAEL003141 PE=4 SV=1
CCG019657.1	1766	32	1031	0.77861354	20.4088241	4.712141821	Up	1.77E-13	5.17E-13	AAEL009987-PA OS=Aedes aegypti GN=AAEL009987 PE=4 SV=1
CCG002971.1	1746	221	1031	5.43889539	20.64260215	1.924239262	Up	0	0	AAEL009704-PA OS=Aedes aegypti GN=AAEL009704 PE=4 SV=1
CCG010251.1	978	2964	1031	130.227294	36.85274371	-1.821187911	Down	0	0	AAEL003453-PA OS=Aedes aegypti GN=AAEL003453 PE=4 SV=1
CCG007922.1	3063	66	1032	0.92589046	11.77830285	3.66914634	Up	0	0	Raptorra OS=Culex quinquefasciatus GN=CpipJ_CPIJ004643 PE=4 SV=1

CCG011478.1	1533	156	1032	4.37265401	23.53355618	2.42813824	Up	1.42E-13	4.26E-13	Serine/threonine-protein phosphatase (Fragment) OS=Aedes aegypti GN=AAEL008930 PE=3 SV=1
CCG000910.1	579	160	1032	11.8741926	62.3090529	2.391612364	Up	4.24E-14	1.45E-13	AAEL011233-FA OS=Aedes aegypti GN=AAEL011233 PE=4 SV=1
CCG025408.1	1503	68	1033	1.94407316	24.02654684	3.627474901	Up	0	0	AAEL015640-FA OS=Aedes aegypti GN=AAEL015640 PE=4 SV=1
CCG014771.1	825	390	1033	20.3129655	43.77199989	1.107607429	Up	6.59E-13	1.68E-12	AAEL012502-FA OS=Aedes aegypti GN=AAEL012502 PE=4 SV=1
CCG009584.1	909	104	1034	4.91622926	39.76552055	3.015893956	Up	0	0	AAEL004976-FA OS=Aedes aegypti GN=AAEL004976 PE=4 SV=1
CCG011307.1	2751	1778	1034	27.7717878	13.13953405	-1.079705935	Down	3.47E-85	1.78E-84	AAEL005439-FA OS=Aedes aegypti GN=AAEL005439 PE=4 SV=1
CCG014290.1	3506	79	1035	0.96822848	10.31997047	3.413947508	Up	0	0	AAEL014377-FA OS=Aedes aegypti GN=AAEL014377 PE=4 SV=1
CCG022025.2	1799	6124	1035	146.273849	20.11218258	-2.862530311	Down	0	0	AAEL005336-FA OS=Aedes aegypti GN=AAEL005336 PE=3 SV=1
CCG013240.1	894	274	1036	13.1696949	40.51093371	1.621089409	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017153 PE=4 SV=1
CCG012564.1	2172	381	1036	7.5375087	16.67438984	1.145474304	Up	0	0	Semaphorin OS=Culex quinquefasciatus GN=CpipJ_CPIJ001595 PE=4 SV=1
CCG016276.1	816	42	1037	2.21167752	44.4261434	4.32819596	Up	7.66E-14	2.47E-13	AAEL000046-FA OS=Aedes aegypti GN=AAEL000046 PE=4 SV=1
CCG021249.1	3504	276	1037	3.38460238	10.34581422	1.611988926	Up	0	0	AAEL008324-FA OS=Aedes aegypti GN=AAEL008324 PE=4 SV=1
CCG010681.1	2193	159	1038	3.11545271	16.54659886	2.409020977	Up	5.53E-14	1.84E-13	AAEL013968-FA OS=Aedes aegypti GN=AAEL013968 PE=4 SV=1
CCG015982.1	1065	301	1038	12.1444978	34.0720106	1.488284255	Up	0	0	AAEL003903-FA OS=Aedes aegypti GN=AAEL003903 PE=4 SV=1
CCG026552.1	792	21	1039	1.13934902	45.86066864	5.33097572	Up	1.76E-10	3.72E-10	AAEL004798-FA OS=Aedes aegypti GN=AAEL004798 PE=4 SV=1

CCG017529.1	1866	82	1039	1.88827344	19.46497833	3.365741138	Up	0	0	AAEL012838-PA OS=Aedes aegypti GN=AAEL012838 PE=4 SV=1
CCG017199.1	3915	105	1039	1.15244499	9.277560553	3.009047625	Up	0	0	AAEL002339-PA OS=Aedes aegypti GN=AAEL002339 PE=4 SV=1
CCG019474.1	2841	138	1039	2.08723104	12.78481153	2.614768686	Up	1.49E-13	4.45E-13	Phosphatase 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ012238 PE=4 SV=1
CCG000292.1	2766	96	1040	1.49135738	13.14410985	3.139718516	Up	0	0	AAEL006031-PA OS=Aedes aegypti GN=AAEL006031 PE=4 SV=1
CCG007222.1	1806	100	1040	2.37927656	20.13101209	3.080824827	Up	0	0	AAEL005074-PA OS=Aedes aegypti GN=AAEL005074 PE=4 SV=1
CCG010389.1	2961	242	1040	3.51187969	12.27848965	1.80581778	Up	3.87E-13	1.03E-12	AAEL001332-PA OS=Aedes aegypti GN=AAEL001332 PE=4 SV=1
CCG000248.1	1452	81	1041	2.39707197	25.06306207	3.386217554	Up	4.46E-14	1.51E-13	AAEL007656-PA OS=Aedes aegypti GN=AAEL007656 PE=4 SV=1
CCG027167.1	2673	270	1041	4.34037723	13.61450285	1.64925196	Up	2.59E-13	7.26E-13	AAEL013903-PB OS=Aedes aegypti GN=AAEL013903 PE=4 SV=1
CCG019118.2	1422	330	1041	9.97187934	25.59181865	1.359745343	Up	1.76E-13	5.16E-13	AAEL002137-PB OS=Aedes aegypti GN=AAEL002137 PE=4 SV=1
CCG014163.1	5542	321	1042	2.48886409	6.572812053	1.401023279	Up	0	0	-
CCG000427.1	1519	167	1043	4.72412487	24.0036094	2.345132354	Up	1.62E-13	4.78E-13	CCA-adding enzyme OS=Culex quinquefasciatus GN=CpipJ_CPIJ013325 PE=3 SV=1
CCG025028.1	717	376	1043	22.5336405	50.85283497	1.174247795	Up	3.71E-14	1.28E-13	AAEL009109-PA OS=Aedes aegypti GN=AAEL009109 PE=4 SV=1
CCG022026.1	1961	2032	1043	44.5254976	18.5933109	-1.25984804	Down	6.69E-124	3.69E-123	AAEL014960-PA (Fragment) OS=Aedes aegypti GN=AAEL014960 PE=4 SV=1
CCG016800.1	1341	269	1044	8.61958136	27.21583963	1.658756838	Up	0	0	AAEL005346-PA OS=Aedes aegypti GN=AAEL005346 PE=4 SV=1
CCG010217.1	2361	194	1045	3.53076176	15.47285016	2.131687589	Up	5.97E-13	1.53E-12	AAEL011862-PA OS=Aedes aegypti GN=AAEL011862 PE=4 SV=1
CCG007403.1	6964	384	1045	2.36938226	5.245749458	1.14663793	Up	1.03E-12	2.55E-12	AAEL004064-PA OS=Aedes aegypti GN=AAEL004064 PE=4 SV=1

CCG013057.1	3705	139	1046	1.61208991	9.869462214	2.614039267	Up	1.46E-13	4.37E-13	r-box and leucine-rich repeat protein 11 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008371 PE=4 SV=1
CCG006878.1	2640	236	1046	3.84123385	13.85089299	1.850337291	Up	4.60E-13	1.20E-12	AAEL007300-PA OS=Aedes aegypti GN=AAEL007360 PE=4 SV=1
CCG025397.1	2189	45	1047	0.88334311	16.72056454	4.242505834	Up	0	0	Chromatin regulatory protein sir2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006725 PE=4 SV=1
CCG011720.1	1578	341	1047	9.28560171	23.19475018	1.320731002	Up	2.98E-13	8.25E-13	AAEL013635-PA OS=Aedes aegypti GN=AAEL013635 PE=4 SV=1
CCG003170.1	2679	394	1047	6.31955037	13.66230526	1.112307111	Up	5.49E-13	1.41E-12	Adenosine A2 receptor OS=Culex quinquefasciatus GN=CpipJ_CPIJ016092 PE=4 SV=1
CCG010631.1	4098	2102	1047	22.0406008	8.931507023	-1.303188023	Down	5.50E-135	3.07E-134	AAEL004319-PA (Fragment) OS=Aedes aegypti GN=AAEL004319 PE=4 SV=1
CCG027646.1	1377	225	1048	7.02119847	26.60586351	1.921955014	Up	0	0	AAEL007313-PA OS=Aedes aegypti GN=AAEL007313 PE=4 SV=1
CCG000592.1	2328	2172	1048	40.0903194	15.73723112	-1.349072182	Down	5.08E-147	2.87E-146	AAEL011393-PB OS=Aedes aegypti GN=AAEL011393 PE=4 SV=1
CCG027226.1	2417	2896	1048	51.4854578	15.15774682	-1.764109682	Down	7.94E-290	4.97E-289	AAEL012888-PA (Fragment) OS=Aedes aegypti GN=AAEL012888 PE=4 SV=1
CCG016499.1	2643	129	1049	2.09727422	13.87485143	2.725884911	Up	0	0	AAEL005152-PA OS=Aedes aegypti GN=AAEL005152 PE=4 SV=1
CCG027694.1	2172	184	1049	3.64016168	16.88362446	2.21355021	Up	3.33E-13	9.08E-13	Zinc finger protein ZTT OS=Culex quinquefasciatus GN=CpipJ_CPIJ000636 PE=4 SV=1
CCG003584.1	639	425	1049	28.5792445	57.38847001	1.005793135	Up	1.58E-12	3.80E-12	Guanylate kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007795 PE=4 SV=1
CCG001842.1	1787	4982	1049	119.795869	20.5211149	-2.54539714	Down	0	0	AAEL008167-PB OS=Aedes aegypti GN=AAEL008167 PE=3 SV=1

CCG006780.1	771	224	1050	12.4840734	47.60854813	1.931131894	Up	2.04E-13	5.88E-13	AAEL007522-PA OS=Aedes aegypti GN=AAEL007522 PE=4 SV=1
CCG009233.1	741	289	1050	16.7587764	49.53601972	1.563561134	Up	0	0	AAEL000649-PA OS=Aedes aegypti GN=AAEL000649 PE=4 SV=1
CCG000092.1	1290	402	1050	13.3905685	28.45441133	1.087435125	Up	0	0	AAEL007280-PA OS=Aedes aegypti GN=AAEL007280 PE=4 SV=1
CCG027844.1	1777	363	1051	8.77772294	20.67594197	1.23603442	Up	0	0	AAEL012973-PA OS=Aedes aegypti GN=AAEL012973 PE=4 SV=1
CCG002100.1	1659	2074	1051	53.7186435	22.14656352	-1.278340021	Down	2.60E-129	1.44E-128	AAEL006150-PA OS=Aedes aegypti GN=AAEL006150 PE=4 SV=1
CCG013625.1	1851	73	1052	1.69464648	19.86823726	3.551407634	Up	0	0	AAEL005858-PA OS=Aedes aegypti GN=AAEL005858 PE=4 SV=1
CCG015406.1	957	312	1052	14.0089417	38.42853413	1.455829974	Up	9.28E-14	2.93E-13	AAEL011753-PA OS=Aedes aegypti GN=AAEL011753 PE=4 SV=1
CCG007519.1	2309	99	1053	1.84235761	15.94242765	3.113246305	Up	0	0	AAEL008519-PA OS=Aedes aegypti GN=AAEL008519 PE=4 SV=1
CCG003938.1	651	167	1053	11.022958	56.54541542	2.358898632	Up	1.62E-13	4.77E-13	AAEL002980-PA OS=Aedes aegypti GN=AAEL002980 PE=4 SV=1
CCG015204.1	854	193	1053	9.71095876	43.10429209	2.150145888	Up	6.08E-14	2.02E-13	-
CCG018468.1	2091	268	1053	5.50735958	17.60452675	1.676513734	Up	0	0	AAEL012020-PA OS=Aedes aegypti GN=AAEL012020 PE=4 SV=1
CCG014136.1	2117	377	1053	7.65214452	17.38831622	1.184182212	Up	4.66E-13	1.21E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003701 PE=4 SV=1
CCG020144.1	933	195	1054	8.9808127	39.49198684	2.136642042	Up	2.26E-13	6.47E-13	AAEL002139-PA OS=Aedes aegypti GN=AAEL002139 PE=4 SV=1
CCG003827.1	1244	375	1054	12.9530952	29.61899013	1.19322557	Up	0	0	AAEL009879-PA OS=Aedes aegypti GN=AAEL009879 PE=4 SV=1
CCG013244.3	2274	6609	1054	124.884334	16.20317666	-2.946243925	Down	0	0	(Z) long form OS=Culex quinquefasciatus GN=CpipJ_CPIJ017150 PE=4 SV=1
CCG011229.1	1347	152	1055	4.84884904	27.38009057	2.497412974	Up	8.39E-14	2.69E-13	AAEL010343-PA (Fragment) OS=Aedes aegypti GN=AAEL010343 PE=4 SV=1

CCG023618.2	3057	311	1055	4.3714712	12.06443637	1.464569717	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008646 PE=4 SV=1
CCG004012.2	1443	32	1056	0.95289779	25.58277219	4.746707323	Up	1.77E-13	5.16E-13	AAEL011358-PA OS=Aedes aegypti GN=AAEL011358 PE=4 SV=1
CCG001642.1	1890	330	1056	7.50265207	19.53224353	1.380385109	Up	1.76E-13	5.16E-13	AAEL013891-PA OS=Aedes aegypti GN=AAEL013891 PE=4 SV=1
CCG017468.1	1697	1913	1056	48.4390703	21.75364777	-1.154913835	Down	7.06E-102	3.75E-101	AGAP000295-PA OS=Anopheles gambiae GN=AgaP_AGAP000295 PE=4 SV=4
CCG018292.1	1748	117	1057	2.87612068	21.13895798	2.877708146	Up	9.44E-14	2.97E-13	AAEL007726-PA OS=Aedes aegypti GN=AAEL007726 PE=4 SV=1
CCG012657.1	1143	129	1057	4.84960259	32.32799523	2.73684561	Up	0	0	AAEL001922-PA OS=Aedes aegypti GN=AAEL001922 PE=4 SV=1
CCG019388.1	2589	143	1057	2.37337661	14.27226672	2.588201528	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ019663 PE=4 SV=1
CCG001598.1	3313	411	1057	5.33068546	11.15330472	1.065078282	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011525 PE=4 SV=1
CCG009259.1	486	17	1058	1.50305656	76.10258606	5.661974275	Up	3.39E-09	6.89E-09	-
CCG016429.1	1782	171	1058	4.12335837	20.75525074	2.331584601	Up	3.42E-13	9.26E-13	AAEL012413-PA OS=Aedes aegypti GN=AAEL012413 PE=4 SV=1
CCG007981.1	1248	8348	1058	287.428962	29.63610323	-3.277777769	Down	0	0	Phosphoglycerate kinase OS=Aedes aegypti GN=PGK PE=3 SV=1
CCG014166.1	2120	64	1059	1.29719954	17.46264863	3.750800078	Up	0	0	Actin OS=Culex quinquefasciatus GN=CpipJ_CPIJ003226 PE=3 SV=1
CCG027004.1	1134	224	1059	8.48784881	32.64622143	1.943445156	Up	2.04E-13	5.87E-13	AAEL010784-PA OS=Aedes aegypti GN=AAEL010784 PE=4 SV=1
CCG016492.2	1050	5035	1059	206.050108	35.25791914	-2.546975985	Down	0	0	AAEL006171-PA (Fragment) OS=Aedes aegypti GN=AAEL006171 PE=4 SV=1

CCG016760.1	2811	102	1061	1.55920062	13.19485295	3.081096803	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_12961 PE=4 SV=1
CCG004023.3	1181	166	1061	6.03977641	31.40620801	2.378482713	Up	1.83E-13	5.31E-13	integrator complex subunit 12 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008030 PE=4 SV=1
CCG024049.1	1626	273	1061	7.21447574	22.81102808	1.660765004	Up	1.53E-14	5.41E-14	AAEL001656-PA OS=Aedes aegypti GN=AAEL001656 PE=4 SV=1
CCG027505.1	727	29	1062	1.71406094	51.06697377	4.896900259	Up	6.66E-13	1.69E-12	AAEL004492-PA OS=Aedes aegypti GN=AAEL004492 PE=4 SV=1
CCG017481.1	1808	84	1062	1.99638148	20.53412054	3.362563832	Up	0	0	AAEL014729-PA (Fragment) OS=Aedes aegypti GN=AAEL014729 PE=4 SV=1
CCG015523.1	1266	2194	1062	74.4672968	29.32518952	-1.344466556	Down	1.03E-147	5.81E-147	AAEL004023-PA OS=Aedes aegypti GN=AAEL004023 PE=4 SV=1
CCG025208.1	2580	31	1063	0.51630301	14.40335202	4.802042775	Up	2.40E-13	6.76E-13	AAEL011811-PA OS=Aedes aegypti GN=AAEL011811 PE=3 SV=1
CCG020098.1	801	121	1063	6.49105854	46.39281924	2.837375848	Up	0	0	-
CCG002368.1	642	180	1063	12.0475891	57.88262961	2.264385989	Up	3.32E-13	9.06E-13	AAEL004121-PA OS=Aedes aegypti GN=AAEL004121 PE=4 SV=1
CCG021707.1	3900	401	1063	4.41817015	9.528371336	1.108780659	Up	1.05E-13	3.28E-13	AAEL006958-PA OS=Aedes aegypti GN=AAEL006958 PE=4 SV=1
CCG021402.1	719	328	1064	19.6023268	51.73241514	1.400043635	Up	1.92E-13	5.53E-13	AAEL005928-PA OS=Aedes aegypti GN=AAEL005928 PE=4 SV=1
CCG021756.1	1293	349	1064	11.5981728	28.7669037	1.310512413	Up	4.55E-14	1.54E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_06292 PE=4 SV=1
CCG000915.1	2267	422	1064	7.99877724	16.40741354	1.036496451	Up	5.52E-13	1.42E-12	AAEL013058-PA OS=Aedes aegypti GN=AAEL013058 PE=4 SV=1
CCG002116.1	1143	24	1066	0.90225165	32.60325725	5.175342426	Up	1.93E-11	4.23E-11	AAEL010862-PA OS=Aedes aegypti GN=AAEL010862 PE=4 SV=1
CCG021456.1	2235	359	1066	6.9020737	16.6736121	1.272464893	Up	0	0	AAEL003519-PA OS=Aedes aegypti GN=AAEL003519 PE=4 SV=1
CCG003020.1	1434	412	1066	12.3455583	25.98711509	1.073804399	Up	3.41E-13	9.25E-13	AAEL010008-PA OS=Aedes aegypti GN=AAEL010008 PE=4 SV=1

CCG026472.1	8151	2057	1066	10.8439141	4.571895846	-1.246021152	Down	2.88E-123	1.59E-122	AAEL011566-PA OS=Aedes aegypti GN=AAEL011566 PE=4 SV=1
CCG006826.1	1546	146	1067	4.05794389	24.12709011	2.571833106	Up	9.33E-15	3.34E-14	AAEL011554-PA OS=Aedes aegypti GN=AAEL011554 PE=4 SV=1
CCG011154.1	3073	94	1068	1.31440125	12.14950849	3.208420284	Up	0	0	Rituper OS=Culex quinquefasciatus GN=CpipJ_CPIJ004355 PE=4 SV=1
CCG024556.1	1937	166	1068	3.68248629	19.27487847	2.387969704	Up	1.83E-13	5.31E-13	AAEL014906-PA (Fragment) OS=Aedes aegypti GN=AAEL014906 PE=4 SV=1
CCG024384.1	3217	186	1068	2.48441736	11.60566975	2.223850324	Up	6.43E-13	1.64E-12	AAEL005480-PA OS=Aedes aegypti GN=AAEL005480 PE=4 SV=1
CCG013203.1	1941	345	1068	7.63758807	19.23515693	1.332556584	Up	9.59E-13	2.38E-12	AGAP007911-PA (Fragment) OS=Anopheles gambiae GN=AGAP007911 PE=4 SV=3
CCG013123.1	1569	129	1069	3.53288449	23.81797187	2.753132086	Up	0	0	AAEL012628-PA OS=Aedes aegypti GN=AAEL012628 PE=4 SV=1
CCG027278.1	561	250	1069	19.1487231	66.61389995	1.798575057	Up	2.10E-13	6.02E-13	AAEL004379-PA OS=Aedes aegypti GN=AAEL004379 PE=4 SV=1
CCG024170.1	2370	70	1071	1.26914828	15.79760102	3.637772952	Up	0	0	AAEL002647-PA OS=Aedes aegypti GN=AAEL002647 PE=4 SV=1
CCG023262.1	1018	142	1071	5.99381367	36.77830493	2.617308849	Up	2.62E-13	7.35E-13	AAEL007167-PA OS=Aedes aegypti GN=AAEL007167 PE=4 SV=1
CCG015738.1	1092	300	1072	11.8048721	34.31801529	1.539583704	Up	1.10E-13	3.42E-13	AAEL011826-PA OS=Aedes aegypti GN=AAEL011826 PE=4 SV=1
CCG013636.1	2106	310	1072	6.32507964	17.79452645	1.492277989	Up	2.14E-13	6.15E-13	AAEL004936-PA OS=Aedes aegypti GN=AAEL004936 PE=4 SV=1
CCG006523.1	1302	94	1074	3.10226963	28.83655089	3.21650263	Up	0	0	AAEL012100-PA OS=Aedes aegypti GN=AAEL012100 PE=4 SV=1
CCG025389.1	1888	152	1074	3.45942779	19.88622312	2.523163968	Up	8.39E-14	2.69E-13	AAEL010001-PA OS=Aedes aegypti GN=AAEL010001 PE=4 SV=1
CCG027269.1	444	193	1074	18.6782855	84.56123706	2.178634445	Up	6.08E-14	2.02E-13	AAEL004121-PA OS=Aedes aegypti GN=AAEL004121 PE=4 SV=1
CCG011437.1	1215	216	1074	7.63906393	30.90139033	2.01620398	Up	1.38E-13	4.17E-13	AAEL000956-PA OS=Aedes aegypti GN=AAEL000956 PE=4 SV=1

CCG005887.1	729	13	1075	0.76626413	51.55027096	6.07199443	Up	6.34E-08	1.23E-07	NOS protein OS=Aedes aegypti GN=nos PE=2 SV=1
CCG004041.1	714	158	1075	9.50870878	52.63325985	2.4686534	Up	0	0	AAEL006473-PA OS=Aedes aegypti GN=AAEL006473 PE=4 SV=1
CCG020465.1	12599	325	1075	1.1084343	2.982788121	1.42813824	Up	1.19E-13	3.65E-13	AAEL014246-PA OS=Aedes aegypti GN=AAEL014248 PE=4 SV=1
CCG017784.1	2562	332	1075	5.56828723	14.66828553	1.397394717	Up	0	0	AAEL012470-PA OS=Aedes aegypti GN=AAEL012470 PE=4 SV=1
CCG014891.1	3710	433	1075	5.01506606	10.12941982	1.014210934	Up	9.43E-13	2.34E-12	AAEL009529-PA OS=Aedes aegypti GN=AAEL009529 PE=4 SV=1
CCG006993.1	715	436	1075	26.2025235	52.5596469	1.004249824	Up	1.61E-13	4.76E-13	AAEL007414-PA OS=Aedes aegypti GN=AAEL007414 PE=4 SV=1
CCG019031.1	1800	166	1076	3.96276441	20.897281	2.398736135	Up	1.83E-13	5.32E-13	AAEL010596-PA OS=Aedes aegypti GN=AAEL010598 PE=4 SV=1
CCG018245.1	1017	403	1076	17.0273383	36.98633806	1.119139538	Up	0	0	AAEL011560-PA OS=Aedes aegypti GN=AAEL011560 PE=4 SV=1
CCG018623.1	1718	30	1077	0.75034461	21.91505476	4.868225143	Up	3.71E-13	9.93E-13	AAEL002036-PA OS=Aedes aegypti GN=AAEL002036 PE=4 SV=1
CCG000982.1	531	118	1077	9.54882991	70.90407549	2.892472689	Up	0	0	AAEL008843-PA OS=Aedes aegypti GN=AAEL008843 PE=4 SV=1
CCG005375.1	1625	188	1077	4.97126776	23.16927021	2.220526887	Up	1.80E-13	5.24E-13	AAEL011995-PA OS=Aedes aegypti GN=AAEL011995 PE=4 SV=1
CCG013470.2	858	4936	1077	247.201177	43.88119357	-2.494010941	Down	0	0	-
CCG007784.1	1125	102	1078	3.8959226	33.49779765	3.104029325	Up	0	0	AAEL014388-PA OS=Aedes aegypti GN=AAEL014388 PE=4 SV=1
CCG020628.1	601	7004	1078	500.765426	62.70386416	-2.997508702	Down	0	0	AAEL005536-PA OS=Aedes aegypti GN=AAEL005538 PE=4 SV=1
CCG027731.1	1224	114	1079	4.00208313	30.81697764	2.944902339	Up	8.48E-14	2.70E-13	AAEL014159-PA OS=Aedes aegypti GN=AAEL014159 PE=4 SV=1
CCG007330.1	2004	232	1079	4.97454013	18.82234563	1.919811358	Up	2.91E-13	8.05E-13	AAEL000004-PA OS=Aedes aegypti GN=AAEL000004 PE=4 SV=1
CCG017450.1	2040	326	1079	6.8667321	18.49018659	1.429064199	Up	6.17E-13	1.57E-12	AAEL013208-PA OS=Aedes aegypti GN=AAEL013208 PE=4 SV=1
CCG011605.1	1384	7678	1079	238.382675	27.25432127	-3.128722492	Down	0	0	AAEL009414-PA OS=Aedes aegypti GN=AAEL009414 PE=4 SV=1

CCG017771.1	1032	86	1080	3.58081122	36.58424313	3.352864046	Up	0	0	Replication factor C subunit 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000302 PE=4 SV=1
CCG021648.1	1582	4624	1080	125.595482	23.86532169	-2.395796882	Down	0	0	AAEL000526-PA OS=Aedes aegypti GN=AAEL000328 PE=4 SV=1
CCG003579.1	798	39	1081	2.10002462	47.35576089	4.495061793	Up	7.33E-14	2.38E-13	-
CCG003423.1	897	126	1081	6.03588245	42.12920534	2.803184088	Up	1.34E-13	4.06E-13	AAEL008849-PA OS=Aedes aegypti GN=AAEL008849 PE=4 SV=1
CCG012549.1	1888	384	1081	8.73960704	20.01583538	1.195501511	Up	1.03E-12	2.55E-12	AAEL002805-PA OS=Aedes aegypti GN=AAEL002805 PE=4 SV=1
CCG023275.1	582	125	1082	9.22889489	64.9911606	2.816013703	Up	2.35E-14	8.22E-14	AAEL013274-PA OS=Aedes aegypti GN=AAEL013274 PE=3 SV=1
CCG000398.2	1435	169	1082	5.06054714	26.3587843	2.380918551	Up	0	0	AAEL008406-PA OS=Aedes aegypti GN=AAEL008408 PE=4 SV=1
CCG008527.1	1797	205	1082	4.90194524	21.04889008	2.102317888	Up	1.83E-13	5.31E-13	AAEL010006-PA OS=Aedes aegypti GN=AAEL010006 PE=4 SV=1
CCG021269.1	2732	370	1082	5.81947357	13.84511547	1.250416527	Up	0	0	AAEL005080-PA OS=Aedes aegypti GN=AAEL005080 PE=4 SV=1
CCG010778.1	1457	29	1083	0.85526582	25.98477265	4.925149736	Up	6.66E-13	1.69E-12	Eukaryotic translation initiation factor 4 gamma 3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009979 PE=4 SV=1
CCG017620.1	5136	275	1084	2.30075487	7.37826558	1.681174437	Up	4.31E-13	1.14E-12	AAEL014900-PA OS=Aedes aegypti GN=AAEL014900 PE=4 SV=1
CCG001756.1	3493	231	1085	2.84168586	10.85878337	1.93404349	Up	1.29E-13	3.97E-13	AAEL004538-PA OS=Aedes aegypti GN=AAEL004538 PE=3 SV=1
CCG012948.1	1647	413	1085	10.7750458	23.02958731	1.09579456	Up	5.88E-13	1.51E-12	AAEL006372-PA OS=Aedes aegypti GN=AAEL006372 PE=4 SV=1
CCG002449.1	1386	111	1086	3.44129909	27.3915502	2.992705725	Up	0	0	AAEL012140-PA OS=Aedes aegypti GN=AAEL012140 PE=4 SV=1
CCG003026.1	3363	316	1086	4.03759623	11.28893505	1.483340843	Up	3.23E-13	8.83E-13	AAEL009096-PA OS=Aedes aegypti GN=AAEL009096 PE=4 SV=1
CCG019947.1	12702	423	1086	1.43097132	2.988874868	1.062607739	Up	1.18E-13	3.65E-13	AAEL012583-PA (Fragment) OS=Aedes aegypti GN=AAEL012583 PE=4 SV=1

CCG008742.1	953	180	1087	8.11600444	39.87371128	2.296596333	Up	3.32E-13	9.05E-13	Ecdysteroid receptor OS=Aedes albopictus GN=EcR PE=2 SV=1
CCG015807.1	2493	431	1087	7.42878284	15.24253785	1.03690537	Up	0	0	Diacylglycerol kinase zeta OS=Culex quinquefasciatus GN=CpipJ_CPIJ018790 PE=4 SV=1
CCG005877.1	2217	175	1088	3.39183742	17.15588865	2.338564933	Up	3.55E-13	9.57E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005943 PE=4 SV=1
CCG018216.1	966	267	1088	11.8767279	39.37329723	1.729080113	Up	5.21E-13	1.35E-12	AAEL001166-PA OS=Aedes aegypti GN=AAEL001166 PE=4 SV=1
CCG013272.2	8127	2395	1088	12.6630386	4.680030162	-1.436033896	Down	5.36E-178	3.13E-177	AAEL010210-PA OS=Aedes aegypti GN=AAEL010210 PE=4 SV=1
CCG022927.1	2667	109	1089	1.75616838	14.27430199	3.022917118	Up	0	0	AAEL001497-PA (Fragment) OS=Aedes aegypti GN=AAEL001497 PE=4 SV=1
CCG005400.1	3663	260	1089	3.04999481	10.3930012	1.76873363	Up	3.22E-13	8.82E-13	AAEL009430-PA OS=Aedes aegypti GN=AAEL009430 PE=4 SV=1
CCG024478.1	999	7	1090	0.30108923	38.14266435	6.985070701	Up	4.31E-06	7.70E-06	Long form D7Bclu1 salivary protein d711 OS=Aedes albopictus PE=2 SV=1
CCG016482.1	2023	95	1090	2.01785704	18.83565086	3.222570015	Up	0	0	AAEL006640-PA OS=Aedes aegypti GN=AAEL006640 PE=3 SV=1
CCG020589.1	1724	204	1090	5.08458577	22.10239077	2.120000282	Up	6.59E-14	2.17E-13	tRNA pseudouridine synthase OS=Aedes aegypti GN=AAEL004663 PE=3 SV=1
CCG019521.1	3789	116	1091	1.31551576	10.06584322	2.935767595	Up	1.16E-13	3.59E-13	AAEL008033-PA OS=Aedes aegypti GN=AAEL008035 PE=4 SV=1
CCG003602.1	3459	240	1091	2.98142131	11.02615784	1.886857995	Up	0	0	AAEL013280-PA OS=Aedes aegypti GN=AAEL013280 PE=4 SV=1
CCG018531.1	1188	278	1091	10.0552073	32.10393936	1.674807517	Up	2.11E-14	7.39E-14	AAEL008687-PA OS=Aedes aegypti GN=AAEL008687 PE=4 SV=1
CCG021139.2	696	326	1091	20.1266286	54.79810339	1.445020436	Up	6.17E-13	1.57E-12	AAEL004480-PA OS=Aedes aegypti GN=AAEL004480 PE=4 SV=1

CCG021902.1	1437	378	1091	11.3031035	26.54104381	1.231506166	Up	1.08E-12	2.65E-12	G-protein signaling modulator OS=Culex quinquefasciatus GN=CpipJ_CPIJ010095 PE=4 SV=1
CCG015167.1	3549	155	1092	1.87667198	10.75639285	2.518945939	Up	3.42E-13	9.28E-13	Structural maintenance of chromosomes protein OS=Aedes aegypti GN=AAEL003449 PE=3 SV=1
CCG000385.1	4929	229	1092	1.99636219	7.744864726	1.955866557	Up	2.46E-13	6.93E-13	AAEL010373-PA OS=Aedes aegypti GN=AAEL010373 PE=4 SV=1
CCG005161.1	1400	272	1093	8.34840558	27.29242608	1.708928048	Up	5.36E-13	1.38E-12	AAEL011058-PA (Fragment) OS=Aedes aegypti GN=AAEL011058 PE=4 SV=1
CCG019869.1	3551	409	1093	4.94920345	10.76017925	1.120433857	Up	3.15E-13	8.65E-13	AAEL003443-PA OS=Aedes aegypti GN=AAEL003445 PE=4 SV=1
CCG013473.1	14730	2921	1093	8.52101798	2.593984828	-1.715855753	Down	4.24E-281	2.65E-280	AAEL003941-PA OS=Aedes aegypti GN=AAEL003941 PE=4 SV=1
CCG005026.1	2367	104	1094	1.88798158	16.15731085	3.097270508	Up	0	0	AAEL013110-PA OS=Aedes aegypti GN=AAEL013110 PE=4 SV=1
CCG024600.1	1281	212	1094	7.11130659	29.8550779	2.069789772	Up	0	0	AAEL005521-PA OS=Aedes aegypti GN=AAEL005521 PE=4 SV=1
CCG016308.1	1044	426	1094	17.5336273	36.63252374	1.063000606	Up	4.37E-13	1.15E-12	AAEL013898-PA OS=Aedes aegypti GN=AAEL013898 PE=4 SV=1
CCG010483.1	954	0	1095	0.001	40.12506611	15.29221615	Up	4.44E-16	1.65E-15	AAEL013300-PA OS=Aedes aegypti GN=AAEL013300 PE=4 SV=1
CCG022679.2	2834	196	1096	2.97179534	13.51950294	2.185635443	Up	7.93E-14	2.55E-13	AAEL008095-PA OS=Aedes aegypti GN=AAEL008095 PE=4 SV=1
CCG013961.1	2067	104	1097	2.16199923	18.55308641	3.101221296	Up	0	0	AAEL010167-PA OS=Aedes aegypti GN=AAEL010167 PE=4 SV=1
CCG014846.1	1663	207	1097	5.34860798	23.06027037	2.108174057	Up	6.88E-15	2.47E-14	AAEL010323-PA OS=Aedes aegypti GN=AAEL010323 PE=4 SV=1
CCG012558.1	2185	256	1097	5.0344403	17.55113484	1.801661014	Up	0	0	AAEL011697-PA OS=Aedes aegypti GN=AAEL011697 PE=3 SV=1

CCG010656.1	1623	176	1099	4.6596878	23.67168587	2.344857256	Up	0	0	Gastrin zinc finger protein XLCGF26.1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006180 PE=4 SV=1
CCG025357.1	1623	234	1099	6.19526673	23.67168587	1.933924155	Up	4.60E-13	1.20E-12	AAEL002162-PA OS=Aedes aegypti GN=AAEL002182 PE=4 SV=1
CCG015795.1	1101	346	1099	13.5036587	34.894774	1.369660647	Up	2.24E-13	6.40E-13	AAEL005646-PA OS=Aedes aegypti GN=AAEL005646 PE=4 SV=1
CCG016798.1	914	134	1100	6.29972039	42.07232434	2.739511822	Up	0	0	AAEL005344-PA OS=Aedes aegypti GN=AAEL005344 PE=4 SV=1
CCG002994.1	1423	5608	1100	169.342426	27.02326384	-2.647669622	Down	0	0	AAEL005630-PA OS=Aedes aegypti GN=AAEL003630 PE=4 SV=1
CCG020807.1	1262	264	1101	8.98891437	30.49846492	1.762517838	Up	0	0	AAEL001166-PA OS=Aedes aegypti GN=AAEL001166 PE=4 SV=1
CCG022012.1	544	230	1102	18.167351	70.81621508	1.962731661	Up	3.85E-13	1.03E-12	Strain nuclear nonnucleoprotein SM D3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005588 PE=4 SV=1
CCG004221.1	1785	345	1102	8.30507476	21.5820846	1.377769161	Up	9.59E-13	2.38E-12	AGAP003794-PA OS=Anopheles gambiae GN=AgaP_AGAP003794 PE=4 SV=5
CCG023948.1	2166	97	1103	1.92431406	17.80192949	3.209617437	Up	3.62E-14	1.25E-13	AAEL006125-PA OS=Aedes aegypti GN=AAEL006125 PE=4 SV=1
CCG024155.1	897	241	1104	11.5448228	43.02557141	1.897948324	Up	2.37E-13	6.72E-13	AGAP011766-PA (Fragment) OS=Anopheles gambiae GN=AGAP011766 PE=3 SV=2
CCG006360.1	2802	421	1105	6.45619496	13.78618695	1.094467435	Up	4.57E-13	1.20E-12	AAEL008714-PA OS=Aedes aegypti GN=AAEL008714 PE=4 SV=1
CCG027196.1	852	22	1106	1.10954714	45.38011046	5.354017255	Up	8.38E-11	1.80E-10	AAEL011906-PA OS=Aedes aegypti GN=AAEL011906 PE=3 SV=1
CCG007336.1	2010	29	1107	0.61996135	19.25314049	4.956771715	Up	6.66E-13	1.69E-12	AAEL007457-PA OS=Aedes aegypti GN=AAEL007457 PE=4 SV=1
CCG003062.1	2978	134	1107	1.93349377	12.99490006	2.74866352	Up	0	0	Protein translocase subunit secA OS=Harpegnathos saltator GN=EAI_00772 PE=4 SV=1

CCG019051.2	1216	2293	1107	81.0276328	31.82468124	-1.348267929	Down	5.86E-155	3.35E-154	AAEL006918-PA OS=Aedes aegypti GN=AAEL006918 PE=4 SV=1
CCG008364.1	1341	2498	1107	80.0435474	28.85817479	-1.471805051	Down	1.56E-192	9.28E-192	Putative night chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ010990 PE=4 SV=1
CCG005191.1	1314	2969	1107	97.0906713	29.45115098	-1.721008667	Down	6.70E-287	4.19E-286	Synaptotagmin I isoform A OS=Culex pipiens GN=sytl PE=2 SV=1
CCG020638.1	1374	216	1108	6.75506745	28.19051722	2.061167868	Up	1.38E-13	4.17E-13	AAEL001412-PA OS=Aedes aegypti GN=AAEL001412 PE=4 SV=1
CCG000450.1	2610	237	1108	3.90184946	14.84052516	1.927312121	Up	2.32E-13	6.62E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012359 PE=4 SV=1
CCG024718.1	4086	2426	1108	25.5126227	9.479630608	-1.428308465	Down	8.09E-179	4.74E-178	AAEL008495-PA OS=Aedes aegypti GN=AAEL008495 PE=4 SV=1
CCG002250.2	3157	227	1109	3.08968317	12.28024357	1.990808367	Up	0	0	AAEL009878-PA OS=Aedes aegypti GN=AAEL009878 PE=4 SV=1
CCG010821.1	2697	307	1109	4.8912527	14.37476045	1.555262009	Up	0	0	AAEL002446-PA (Fragment) OS=Aedes aegypti GN=AAEL002446 PE=4 SV=1
CCG013885.1	2168	124	1110	2.45767855	17.89837971	2.864460855	Up	1.61E-13	4.76E-13	AAEL002098-PA (Fragment) OS=Aedes aegypti GN=AAEL002098 PE=4 SV=1
CCG018830.1	2115	291	1110	5.91214788	18.34689703	1.633781822	Up	6.00E-14	2.00E-13	AAEL000884-PA OS=Aedes aegypti GN=AAEL000884 PE=4 SV=1
CCG023490.1	2376	1978	1110	35.7719424	16.33151819	-1.131169546	Down	7.37E-102	3.91E-101	AAEL000485-PA OS=Aedes aegypti GN=AAEL000485 PE=4 SV=1
CCG009952.2	5464	4277	1110	33.6349844	7.10169971	-2.243726327	Down	0	0	AAEL012587-PA OS=Aedes aegypti GN=AAEL012587 PE=4 SV=1
CCG016440.1	2541	73	1111	1.23447093	15.28478768	3.630131746	Up	0	0	AAEL004368-PA OS=Aedes aegypti GN=AAEL004368 PE=4 SV=1
CCG016663.1	2319	159	1111	2.94617844	16.74801444	2.50707335	Up	5.53E-14	1.84E-13	AAEL000439-PA OS=Aedes aegypti GN=AAEL000439 PE=4 SV=1
CCG020635.1	4341	260	1111	2.57363073	8.946935152	1.797588492	Up	3.22E-13	8.81E-13	AAEL001397-PA OS=Aedes aegypti GN=AAEL001397 PE=3 SV=1

CCG026475.1	2010	341	1111	7.2898903	19.3227092	1.406328376	Up	2.98E-13	8.25E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002173 PE=4 SV=1
CCG006511.1	2810	104	1112	1.59033893	13.83402269	3.120814558	Up	0	0	AAEL009242-PA OS=Aedes aegypti GN=AAEL009242 PE=4 SV=1
CCG024251.1	1816	169	1112	3.99883543	21.40616948	2.42037484	Up	0	0	AAEL003771-PA OS=Aedes aegypti GN=AAEL003771 PE=4 SV=1
CCG024868.1	6414	417	1112	2.79363569	6.060742714	1.117350703	Up	0	0	AAEL013079-PA (Fragment) OS=Aedes aegypti GN=AAEL013079 PE=4 SV=1
CCG024218.1	2370	155	1113	2.8102569	16.41711479	2.546426676	Up	3.42E-13	9.26E-13	AAEL000003-PA OS=Aedes aegypti GN=AAEL000003 PE=4 SV=1
CCG017477.1	2876	313	1113	4.67646973	13.52870725	1.532532234	Up	0	0	Chloride channel protein 7 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008618 PE=4 SV=1
CCG000907.1	956	401	1113	18.0239159	40.69933269	1.175092655	Up	1.05E-13	3.28E-13	AAEL004854-PA OS=Aedes aegypti GN=AAEL004854 PE=4 SV=1
CCG004820.1	1404	146	1114	4.46836272	27.73755009	2.634022162	Up	9.33E-15	3.34E-14	AGAP002348-PA OS=Anopheles gambiae GN=AgaP_AGAP002348 PE=4 SV=5
CCG001395.1	1048	189	1114	7.74931283	37.15984764	2.261604297	Up	2.34E-13	6.65E-13	AAEL008700-PA OS=Aedes aegypti GN=AAEL008700 PE=4 SV=1
CCG007645.1	2176	260	1115	5.13425138	17.91290377	1.802773386	Up	3.22E-13	8.81E-13	AAEL001282-PA OS=Aedes aegypti GN=AAEL001282 PE=4 SV=1
CCG005429.3	906	231	1116	10.9558595	43.06118861	1.974685474	Up	1.29E-13	3.97E-13	AAEL000042-PA OS=Aedes aegypti GN=AAEL000042 PE=4 SV=1
CCG007422.1	966	3951	1116	175.748883	40.38658062	-2.121567615	Down	0	0	AAEL004118-PA OS=Aedes aegypti GN=AAEL004118 PE=4 SV=1
CCG008997.1	1881	295	1117	6.73900676	20.75938073	1.62315553	Up	2.32E-13	6.62E-13	AAEL003248-PA OS=Aedes aegypti GN=AAEL003248 PE=4 SV=1
CCG002881.1	666	421	1117	27.16255	58.63122396	1.110050251	Up	4.57E-13	1.20E-12	AAEL009713-PA OS=Aedes aegypti GN=AAEL009713 PE=4 SV=1
CCG004762.2	1742	205	1118	5.05671389	22.43590897	2.149537577	Up	1.83E-13	5.30E-13	AGAP002258-PB OS=Anopheles gambiae GN=AgaP_AGAP002258 PE=4 SV=1

CCG021278.1	1530	207	1118	5.81355233	25.54467545	2.135530719	Up	6.88E-15	2.47E-14	AAEL010665-PA OS=Aedes aegypti GN=AAEL010665 PE=4 SV=1
CCG026152.1	1177	272	1118	9.93013408	33.20590776	1.741554835	Up	5.36E-13	1.38E-12	AAEL003864-PA OS=Aedes aegypti GN=AAEL003864 PE=4 SV=1
CCG019634.1	1482	223	1119	6.46575629	26.39562194	2.029407625	Up	0	0	AAEL014037-PA OS=Aedes aegypti GN=AAEL014037 PE=4 SV=1
CCG009599.1	2886	321	1119	4.77937797	13.55450856	1.503878038	Up	0	0	AAEL010577-PA OS=Aedes aegypti GN=AAEL010577 PE=4 SV=1
CCG022821.1	1072	22	1120	0.88184157	36.52357275	5.372164602	Up	8.38E-11	1.80E-10	-
CCG022066.1	758	38	1120	2.15415556	51.65339048	4.583668707	Up	4.26E-14	1.45E-13	AAEL004624-PA OS=Aedes aegypti GN=AAEL004624 PE=4 SV=1
CCG016912.1	1095	64	1120	2.51147307	35.75641095	3.831596221	Up	0	0	AAEL011738-PA OS=Aedes aegypti GN=AAEL011738 PE=4 SV=1
CCG025859.1	2735	224	1121	3.51927625	14.32841984	2.025528845	Up	2.04E-13	5.88E-13	AAEL000707-PA OS=Aedes aegypti GN=AAEL000707 PE=4 SV=1
CCG028528.2	2128	299	1121	6.03757079	18.4155208	1.608882092	Up	4.80E-13	1.25E-12	AAEL014389-PA OS=Aedes aegypti GN=AAEL014389 PE=4 SV=1
CCG015846.1	1938	424	1121	9.40101521	20.22096402	1.104963312	Up	0	0	AAEL005009-PA OS=Aedes aegypti GN=AAEL005009 PE=4 SV=1
CCG019595.1	1401	115	1122	3.52713739	27.99656427	2.988680113	Up	6.13E-14	2.03E-13	AAEL007356-PA OS=Aedes aegypti GN=AAEL007356 PE=4 SV=1
CCG013437.1	1282	288	1122	9.65310731	30.59530931	1.664245163	Up	0	0	AAEL003491-PA OS=Aedes aegypti GN=AAEL003491 PE=3 SV=1
CCG026088.1	3795	389	1122	4.40453933	10.33549052	1.230543819	Up	0	0	AAEL012508-PA OS=Aedes aegypti GN=AAEL012508 PE=4 SV=1
CCG009002.1	2904	303	1123	4.48341239	13.51864491	1.592281433	Up	1.37E-13	4.14E-13	AAEL004477-PA OS=Aedes aegypti GN=AAEL004477 PE=3 SV=1
CCG017953.2	5029	141	1124	1.20475891	7.813303458	2.697188172	Up	0	0	AAEL007468-PA OS=Aedes aegypti GN=AAEL007468 PE=4 SV=1
CCG017911.1	1630	144	1124	3.79609925	24.10619822	2.666814523	Up	7.53E-14	2.44E-13	AAEL004128-PA OS=Aedes aegypti GN=AAEL004128 PE=4 SV=1
CCG001790.1	3885	136	1125	1.50421722	10.12305312	2.750559649	Up	1.70E-13	4.99E-13	AAEL002465-PA (Fragment) OS=Aedes aegypti GN=AAEL002465 PE=4 SV=1

CCG018560.1	3088	201	1125	2.79692897	12.73577117	2.186970799	Up	0	0	AAEL002238-PA OS=Aedes aegypti GN=AAEL002238 PE=4 SV=1
CCG002261.1	1367	44	1126	1.38307851	28.79518628	4.379872697	Up	3.91E-14	1.34E-13	AAEL007554-PA OS=Aedes aegypti GN=AAEL007554 PE=4 SV=1
CCG000978.1	1350	121	1126	3.8513614	29.15779233	2.920441079	Up	0	0	AAEL012414-PA OS=Aedes aegypti GN=AAEL012414 PE=4 SV=1
CCG011087.1	414	244	1126	25.3251576	95.0797576	1.908566978	Up	2.78E-13	7.73E-13	AAEL012840-PA OS=Aedes aegypti GN=AAEL012840 PE=4 SV=1
CCG020834.1	6410	2506	1126	16.7990881	6.1408767	-1.451866383	Down	3.05E-189	1.81E-188	-
CCG004390.2	1992	384	1127	8.28332233	19.77810137	1.255622503	Up	1.03E-12	2.55E-12	AAEL009686-PA (Fragment) OS=Aedes aegypti GN=AAEL009686 PE=4 SV=1
CCG014167.1	1777	395	1127	9.5515167	22.17106242	1.214876161	Up	0	0	AAEL002836-PA (Fragment) OS=Aedes aegypti GN=AAEL002836 PE=4 SV=1
CCG011474.1	14145	3315	1127	10.0703196	2.785293596	-1.854208151	Down	0	0	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_05561 PE=3 SV=1
CCG014331.1	1486	51	1128	1.47473517	26.53629623	4.169439214	Up	0	0	AAEL013891-PA OS=Aedes aegypti GN=AAEL013891 PE=4 SV=1
CCG014840.1	1700	84	1128	2.12321042	23.19584482	3.449547133	Up	0	0	AAEL011701-PA OS=Aedes aegypti GN=AAEL011701 PE=4 SV=1
CCG006294.1	1494	353	1129	10.1528222	26.41760005	1.379618601	Up	0	0	AAEL011551-PA OS=Aedes aegypti GN=AAEL011551 PE=4 SV=1
CCG017051.1	1671	369	1129	9.48882829	23.61932644	1.315665969	Up	0	0	AAEL008819-PA OS=Aedes aegypti GN=AAEL008819 PE=4 SV=1
CCG010536.1	303	3154	1129	447.28232	130.2570775	-1.77982397	Down	0	0	AAEL013960-PA OS=Aedes aegypti GN=AAEL013960 PE=4 SV=1
CCG014316.1	264	3812	1129	620.456926	149.4996003	-2.053189429	Down	0	0	AAEL009214-PA OS=Aedes aegypti GN=AAEL009214 PE=4 SV=1
CCG016274.1	1129	139	1130	5.29033934	34.9892407	2.725479188	Up	1.46E-13	4.37E-13	AAEL000076-PA OS=Aedes aegypti GN=AAEL000076 PE=4 SV=1
CCG011060.1	1461	2641	1130	77.6749275	27.03822913	-1.522448325	Down	6.40E-214	3.85E-213	AAEL007452-PA OS=Aedes aegypti GN=AAEL007452 PE=4 SV=1

CCG025840.1	2673	186	1131	2.99003765	14.79154921	2.306537607	Up	6.43E-13	1.64E-12	AAEL007409-PA (Fragment) OS=Aedes aegypti GN=AAEL007409 PE=4 SV=1
CCG004988.1	1761	225	1131	5.49017052	22.45190859	2.031915227	Up	0	0	Apyrase OS=Culex quinquefasciatus GN=CpipJ_CPIJ011013 PE=3 SV=1
CCG004938.1	1221	160	1132	5.63075965	32.41013047	2.525043352	Up	4.24E-14	1.45E-13	AAEL012942-PA OS=Aedes aegypti GN=AAEL012942 PE=4 SV=1
CCG004519.1	1461	193	1132	5.67635782	27.0860844	2.254514409	Up	6.08E-14	2.02E-13	AAEL001691-PA OS=Aedes aegypti GN=AAEL001691 PE=4 SV=1
CCG013315.1	801	377	1132	20.2242072	49.40420637	1.288550733	Up	4.66E-13	1.21E-12	AAEL007179-PA OS=Aedes aegypti GN=AAEL007179 PE=4 SV=1
CCG002957.1	7715	434	1132	2.41722162	5.129328491	1.085420214	Up	0	0	AAEL007931-PA OS=Aedes aegypti GN=AAEL007931 PE=4 SV=1
CCG000094.1	1089	18	1133	0.71024355	36.3707324	5.678320348	Up	1.62E-09	3.33E-09	Long form D7Bclu1 salivary protein d711 OS=Aedes albopictus PE=2 SV=1
CCG013575.1	6984	4887	1133	30.0677396	5.671209562	-2.40648804	Down	0	0	AAEL002228-PA OS=Aedes aegypti GN=AAEL002228 PE=4 SV=1
CCG018583.1	1381	91	1134	2.8314597	28.70578266	3.341723489	Up	0	0	AAEL001324-PA OS=Aedes aegypti GN=AAEL001324 PE=4 SV=1
CCG001940.1	2184	366	1134	7.20097201	18.15141294	1.33381829	Up	0	0	AAEL002488-PA OS=Aedes aegypti GN=AAEL002488 PE=4 SV=1
CCG004254.1	1404	166	1136	5.0804672	28.28532935	2.477020892	Up	1.83E-13	5.32E-13	AAEL007554-PA OS=Aedes aegypti GN=AAEL007554 PE=4 SV=1
CCG004145.1	3900	218	1136	2.40189799	10.18271857	2.083875999	Up	0	0	AAEL007563-PA (Fragment) OS=Aedes aegypti GN=DUOX PE=4 SV=2
CCG000776.1	1689	105	1137	2.67129789	23.53319165	3.139084225	Up	0	0	AAEL000550-PA OS=Aedes aegypti GN=AAEL000550 PE=4 SV=1
CCG019716.1	1776	176	1137	4.25826199	22.38038327	2.393898124	Up	0	0	AAEL003304-PA OS=Aedes aegypti GN=AAEL003304 PE=4 SV=1
CCG009981.1	1431	233	1137	6.99646972	27.77607316	1.989143598	Up	0	0	AAEL001187-PA OS=Aedes aegypti GN=AAEL001187 PE=4 SV=1

CCG023493.1	453	391	1137	37.0886672	87.74295958	1.242304945	Up	9.02E-13	2.25E-12	Putative salivary secreted peptide OS=Aedes albopictus PE=2 SV=1
CCG008099.1	2130	1968	1137	39.701614	18.66082662	-1.089184763	Down	2.05E-95	1.08E-94	AAEL011369-FA OS=Aedes aegypti GN=AAEL011369 PE=4 SV=1
CCG011132.1	1965	83	1138	1.8150066	20.24555673	3.479558615	Up	1.32E-13	4.01E-13	AAEL015340-FA OS=Aedes aegypti GN=AAEL015340 PE=4 SV=1
CCG004813.1	966	125	1138	5.56026587	41.18273185	2.888813761	Up	2.35E-14	8.22E-14	AAEL008833-FA OS=Aedes aegypti GN=AAEL008833 PE=4 SV=1
CCG014171.1	333	4337	1138	559.638856	119.4670239	-2.227883682	Down	0	0	AAEL002825-FA OS=Aedes aegypti GN=AAEL002825 PE=4 SV=1
CCG025551.2	2014	325	1139	6.93404357	19.7703462	1.511569328	Up	1.19E-13	3.66E-13	AAEL008005-FA OS=Aedes aegypti GN=AAEL008005 PE=4 SV=1
CCG017278.1	2126	2093	1139	42.3027538	18.72882279	-1.175491361	Down	1.96E-114	1.06E-113	AAEL000080-FA OS=Aedes aegypti GN=AAEL000080 PE=3 SV=1
CCG010261.1	1164	301	1140	11.1115894	34.23748756	1.623511636	Up	0	0	AAEL002890-FA OS=Aedes aegypti GN=AAEL002890 PE=4 SV=1
CCG023335.1	3180	347	1140	4.68883582	12.53221243	1.41833946	Up	0	0	AAEL002925-FA OS=Aedes aegypti GN=AAEL002925 PE=4 SV=1
CCG007434.1	936	356	1140	16.3431897	42.57738838	1.381397882	Up	7.93E-13	1.99E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_19522 PE=4 SV=1
CCG013059.1	7158	10543	1140	63.2900129	5.567537793	-3.506866509	Down	0	0	AAEL008160-FA OS=Aedes aegypti GN=AAEL008160 PE=4 SV=1
CCG010748.1	1692	0	1141	0.001	23.57410981	14.52491567	Up	4.44E-16	1.64E-15	Putative salivary secreted protein 62k-2 OS=Aedes albopictus PE=2 SV=1
CCG006185.1	1561	116	1141	3.19313851	25.55246239	3.000415285	Up	1.16E-13	3.59E-13	AAEL014000-FA OS=Aedes aegypti GN=AAEL014000 PE=4 SV=1
CCG023678.1	1577	121	1141	3.29698027	25.29321103	2.939533043	Up	0	0	AAEL006400-FA OS=Aedes aegypti GN=AAEL006400 PE=4 SV=1
CCG015618.1	1871	127	1141	2.91670566	21.31875671	2.869711593	Up	1.57E-13	4.67E-13	AAEL007858-FA OS=Aedes aegypti GN=AAEL007858 PE=4 SV=1
CCG002154.1	985	68	1142	2.96643853	40.53030667	3.772197298	Up	0	0	AAEL014289-FA OS=Aedes aegypti GN=AAEL014289 PE=4 SV=1

CCG024053.2	2950	96	1142	1.39833713	13.5330007	3.274697638	Up	0	0	AAEL013624-PA OS=Aedes aegypti GN=AAEL013624 PE=4 SV=1
CCG010370.1	2685	136	1142	2.17649308	14.86865999	2.772197298	Up	1.70E-13	4.98E-13	AAEL001746-PA OS=Aedes aegypti GN=AAEL001746 PE=4 SV=1
CCG007173.1	1577	111	1143	3.02450256	25.3375462	3.066507026	Up	0	0	AAEL000547-PA OS=Aedes aegypti GN=AAEL008547 PE=4 SV=1
CCG019140.1	4651	185	1143	1.70918102	8.591122415	2.329541432	Up	0	0	AAEL002312-PA OS=Aedes aegypti GN=AAEL002312 PE=4 SV=1
CCG002349.1	549	445	1143	34.8297485	72.78198607	1.063261366	Up	0	0	Growth factor receptor-bound protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000896 PE=4 SV=1
CCG012454.1	2982	257	1144	3.70329369	13.41122355	1.856559991	Up	4.14E-13	1.10E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004892 PE=4 SV=1
CCG020537.1	1938	329	1144	7.29465567	20.63584553	1.500240767	Up	0	0	AAEL012355-PA (Fragment) OS=Aedes aegypti GN=AAEL012355 PE=4 SV=1
CCG024779.1	1896	440	1144	9.97187934	21.09296869	1.080824827	Up	0	0	AAEL014130-PA OS=Aedes aegypti GN=AAEL014130 PE=4 SV=1
CCG015344.1	1096	12307	1144	482.507777	36.48929619	-3.725006967	Down	0	0	AAEL000445-PA OS=Aedes aegypti GN=AAEL000445 PE=4 SV=1
CCG022157.1	5478	203	1145	1.59234321	7.30690524	2.19810917	Up	9.28E-14	2.93E-13	AAEL013557-PA OS=Aedes aegypti GN=AAEL013557 PE=4 SV=1
CCG028347.1	2416	370	1145	6.58062989	16.5675608	1.332063626	Up	0	0	-
CCG026500.1	1853	320	1147	7.42056939	21.63904126	1.544034785	Up	0	0	AAEL010065-PA OS=Aedes aegypti GN=AAEL010065 PE=2 SV=1
CCG007913.2	2979	145	1148	2.09151108	13.47166893	2.68731104	Up	0	0	AAEL003540-PA OS=Aedes aegypti GN=AAEL003540 PE=4 SV=1
CCG012873.1	834	1	1149	0.05152246	48.16194246	9.868476286	Up	0.000587	0.00091655	AAEL014128-PA OS=Aedes aegypti GN=AAEL014128 PE=4 SV=1
CCG010700.1	2639	145	1149	2.36097443	15.22056082	2.688567196	Up	0	0	AGAP002097-PA OS=Anopheles gambiae GN=AgaP_AGAP002097 PE=4 SV=5

CCG013501.1	1351	61	1150	1.94015826	29.75723041	3.938994012	Up	1.75E-14	6.17E-14	Uncharacterized protein OS=Anopheles darlingi GN=AND_00721 PE=4 SV=1
CCG022558.1	1764	256	1150	6.23597056	22.7902598	1.86973135	Up	0	0	AAEL003987-PA OS=Aedes aegypti GN=AAEL003987 PE=4 SV=1
CCG026958.2	2490	459	1150	7.92092698	16.14538887	1.027381006	Up	5.70E-13	1.47E-12	AAEL005077-PA OS=Aedes aegypti GN=AAEL005077 PE=4 SV=1
CCG026665.1	1465	55	1151	1.61319823	27.46551301	4.089625608	Up	0	0	AAEL010412-PA OS=Aedes aegypti GN=AAEL010412 PE=4 SV=1
CCG008541.1	1569	343	1151	9.39363861	25.64498188	1.448920556	Up	0	0	AAEL007738-PA OS=Aedes aegypti GN=AAEL007738 PE=4 SV=1
CCG023167.1	1078	426	1151	16.9806187	37.32558123	1.136275702	Up	4.37E-13	1.15E-12	-
CCG019296.1	1633	21	1153	0.55258079	24.68272696	5.481172579	Up	1.76E-10	3.73E-10	AAEL007496-PA OS=Aedes aegypti GN=AAEL007496 PE=4 SV=1
CCG021449.1	1068	89	1153	3.58081122	37.74053663	3.397756571	Up	0	0	AAEL003522-PA OS=Aedes aegypti GN=AAEL003522 PE=4 SV=1
CCG025691.1	1591	300	1153	8.10240125	25.33431371	1.644671311	Up	1.10E-13	3.42E-13	AAEL012330-PA OS=Aedes aegypti GN=AAEL012330 PE=3 SV=1
CCG002461.1	1281	326	1154	10.9353111	31.49246791	1.526012558	Up	6.17E-13	1.58E-12	AAEL013965-PA (Fragment) OS=Aedes aegypti GN=AAEL013965 PE=4 SV=1
CCG025388.1	1659	19	1155	0.49211872	24.33804079	5.628062827	Up	7.73E-10	1.60E-09	AAEL010010-PA OS=Aedes aegypti GN=AAEL010010 PE=4 SV=1
CCG008312.1	3586	245	1155	2.93574595	11.25956767	1.939352401	Up	3.40E-13	9.24E-13	AAEL003573-PA OS=Aedes aegypti GN=AAEL003573 PE=4 SV=1
CCG009134.1	2001	261	1155	5.60474799	20.17831568	1.848084344	Up	8.02E-14	2.58E-13	AAEL005636-PA OS=Aedes aegypti GN=AAEL005636 PE=4 SV=1
CCG013942.1	1869	442	1155	10.1619169	21.60342947	1.088087781	Up	0	0	AAEL006525-PA OS=Aedes aegypti GN=AAEL006525 PE=4 SV=1
CCG021250.1	1914	306	1156	6.86976948	21.11377636	1.619851044	Up	0	0	AAEL008319-PA OS=Aedes aegypti GN=AAEL008319 PE=4 SV=1
CCG012565.1	2076	408	1156	8.44491894	19.46616953	1.204813544	Up	1.01E-12	2.49E-12	AAEL002651-PA OS=Aedes aegypti GN=AAEL002651 PE=4 SV=1
CCG006176.2	930	3776	1156	174.466363	43.45351392	-2.005404163	Down	0	0	AAEL000757-PA OS=Aedes aegypti GN=AAEL000757 PE=4 SV=1

CCG026717.1	1750	308	1157	7.56267329	23.11241499	1.611699812	Up	0	0	AAEL004315-PA OS=Aedes aegypti GN=AAEL004315 PE=4 SV=1
CCG024703.1	2027	316	1157	6.69878448	19.95398432	1.574705605	Up	3.23E-13	8.83E-13	AAEL005552-PA OS=Aedes aegypti GN=AAEL005552 PE=3 SV=1
CCG001285.1	1491	165	1158	4.75520202	27.15069383	2.513410528	Up	1.31E-14	4.67E-14	AAEL005798-PA OS=Aedes aegypti GN=VATB PE=2 SV=1
CCG006823.1	1848	283	1159	6.58032191	21.92459025	1.736319812	Up	0	0	AAEL005461-PA (Fragment) OS=Aedes aegypti GN=AAEL005461 PE=4 SV=1
CCG009311.1	2433	53	1160	0.93604436	16.66732472	4.154301839	Up	0	0	AAEL014341-PA OS=Aedes aegypti GN=AAEL014341 PE=4 SV=1
CCG012268.1	1434	146	1161	4.37488232	28.30303998	2.693640902	Up	9.33E-15	3.34E-14	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011278 PE=4 SV=1
CCG000196.1	3624	231	1161	2.73896487	11.19938171	2.031716419	Up	1.29E-13	3.97E-13	AAEL006139-PA OS=Aedes aegypti GN=AAEL006139 PE=3 SV=1
CCG020157.1	2853	84	1162	1.26514466	14.23817652	3.492390134	Up	0	0	AGAP005525-PA OS=Anopheles gambiae GN=AGAP005525 PE=4 SV=1
CCG007039.1	414	138	1162	14.3232449	98.11960775	2.7761831	Up	1.49E-13	4.45E-13	-
CCG000390.1	1479	182	1162	5.28768878	27.46552915	2.376912917	Up	0	0	AAEL010378-PA OS=Aedes aegypti GN=AAEL010378 PE=4 SV=1
CCG012391.1	1641	433	1162	11.3381445	24.75412408	1.126484342	Up	9.43E-13	2.34E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000417 PE=4 SV=1
CCG006081.1	1572	0	1163	0.001	25.86289815	14.65859633	Up	4.44E-16	1.65E-15	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG013715.1	636	177	1163	11.9585582	63.92527655	2.418343035	Up	0	0	AAEL001795-PA OS=Aedes aegypti GN=AAEL001795 PE=3 SV=1
CCG023621.1	528	145	1164	11.8004006	77.06711016	2.707279457	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005778 PE=4 SV=1
CCG015072.1	2049	186	1164	3.90062013	19.85916748	2.348029736	Up	6.43E-13	1.64E-12	AAEL004708-PA (Fragment) OS=Aedes aegypti GN=AAEL004708 PE=4 SV=1

CCG011901.1	3903	233	1164	2.56519297	10.42568131	2.023002402	Up	0	0	AAEL001387-PA OS=Aedes aegypti GN=AAEL001387 PE=4 SV=1
CCG021336.1	1026	340	1164	14.2394832	39.66026722	1.477797611	Up	8.86E-14	2.80E-13	AAEL013394-PA OS=Aedes aegypti GN=AAEL013394 PE=3 SV=1
CCG021416.1	12680	384	1164	1.30129165	3.209103641	1.302226046	Up	1.03E-12	2.55E-12	AAEL010974-PA OS=Aedes aegypti GN=AAEL010974 PE=4 SV=1
CCG000279.1	918	446	1164	20.8763634	44.32618101	1.086288647	Up	0	0	AAEL011803-PA OS=Aedes aegypti GN=AAEL011803 PE=4 SV=1
CCG016002.1	3948	165	1165	1.79584757	10.31570224	2.522105229	Up	1.31E-14	4.67E-14	AAEL002339-PA OS=Aedes aegypti GN=AAEL002339 PE=4 SV=1
CCG020900.1	5208	334	1165	2.75573951	7.819967826	1.504723151	Up	3.15E-13	8.65E-13	AAEL013534-PA (Fragment) OS=Aedes aegypti GN=AAEL013534 PE=4 SV=1
CCG003554.1	2385	349	1166	6.28781441	17.09071309	1.442582051	Up	4.55E-14	1.54E-13	AAEL012247-PA OS=Aedes aegypti GN=AAEL012247 PE=4 SV=1
CCG021410.1	2871	436	1166	6.52553267	14.19761432	1.121480952	Up	1.61E-13	4.76E-13	AAEL013386-PA OS=Aedes aegypti GN=AAEL013386 PE=4 SV=1
CCG012636.1	1140	448	1167	16.8863518	35.78623596	1.083547127	Up	0	0	Membrann OS=Culex quinquefasciatus GN=CpipJ_CPIJ004746 PE=4 SV=1
CCG007679.1	4731	2237	1167	20.3177545	8.623189388	-1.236447492	Down	2.81E-132	1.56E-131	AAEL004098-PA OS=Aedes aegypti GN=AAEL004098 PE=4 SV=1
CCG027138.1	17690	2149	1169	5.22000903	2.310131461	-1.176077349	Down	1.55E-117	8.43E-117	AAEL005179-PA OS=Aedes aegypti GN=AAEL005179 PE=4 SV=1
CCG018589.1	1230	8691	1169	303.617857	33.22457362	-3.191934052	Down	0	0	Alginate kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007538 PE=3 SV=1
CCG023441.1	1614	375	1170	9.9836744	25.34150175	1.343859233	Up	0	0	AAEL002658-PA OS=Aedes aegypti GN=AAEL002658 PE=4 SV=1
CCG012799.1	1798	3307	1170	79.0327655	22.74815563	-1.796701312	Down	0	0	AAEL005430-PA OS=Aedes aegypti GN=AAEL005430 PE=4 SV=1
CCG003410.1	1812	3655	1170	86.6746027	22.57239725	-1.941049673	Down	0	0	AAEL004439-PA OS=Aedes aegypti GN=AAEL004439 PE=4 SV=1
CCG017547.1	2720	138	1171	2.18008212	15.05005224	2.787314108	Up	1.49E-13	4.45E-13	AAEL009479-PA OS=Aedes aegypti GN=AAEL009479 PE=4 SV=1

CCG019105.1	33052	12110	1171	15.7437821	1.23853752	-3.66807268	Down	0	0	AAEL004204-PA OS=Aedes aegypti GN=AAEL004264 PE=4 SV=1
CCG028574.1	981	118	1172	5.16863271	41.76462832	3.014427009	Up	0	0	AAEL011385-PA OS=Aedes aegypti GN=AAEL011385 PE=4 SV=1
CCG002010.1	3516	174	1172	2.12648857	11.65275892	2.454126562	Up	1.78E-13	5.20E-13	AAEL014631-PA OS=Aedes aegypti GN=AAEL014631 PE=4 SV=1
CCG000879.1	954	237	1172	10.6748712	42.9466461	2.008326809	Up	2.32E-13	6.62E-13	AAEL009765-PA OS=Aedes aegypti GN=AAEL009765 PE=4 SV=1
CCG025184.2	2976	262	1172	3.78295379	13.76717083	1.863647057	Up	0	0	Viterbigenin convervase OS=Culex quinquefasciatus GN=CpipJ_CPIJ011701 PE=4 SV=1
CCG007517.1	414	2607	1172	270.584778	98.96401057	-1.451104809	Down	1.24E-196	7.40E-196	AAEL002587-PA OS=Aedes aegypti GN=AAEL002587 PE=4 SV=1
CCG020269.1	867	23	1173	1.13991222	47.2964921	5.374738546	Up	4.01E-11	8.68E-11	AAEL005325-PA OS=Aedes aegypti GN=AAEL005325 PE=4 SV=1
CCG027539.1	19727	413	1173	0.89960462	2.078676872	1.20830253	Up	5.88E-13	1.51E-12	AAEL011567-PA OS=Aedes aegypti GN=AAEL011567 PE=3 SV=1
CCG024028.1	22197	3628	1173	7.02321022	1.847369404	-1.926658239	Down	0	0	AAEL010061-PA OS=Aedes aegypti GN=AAEL010061 PE=4 SV=1
CCG014428.1	660	144	1174	9.37521482	62.18335899	2.729604895	Up	7.53E-14	2.44E-13	GATA transcription factor GATAb-2 OS=Aedes aegypti PE=2 SV=1
CCG006562.1	2714	173	1174	2.73904351	15.12196644	2.464901669	Up	3.60E-13	9.68E-13	AAEL000157-PA OS=Aedes aegypti GN=AAEL000157 PE=4 SV=1
CCG022989.1	11652	387	1174	1.42716163	3.522229397	1.303340141	Up	1.31E-12	3.18E-12	AGAP005300-PA OS=Anopheles gambiae GN=AGAP005300 PE=2 SV=1
CCG016706.1	628	33	1175	2.25796376	65.40760383	4.856364126	Up	1.31E-13	4.01E-13	-
CCG012299.1	1545	120	1175	3.33745512	26.58639172	2.99386765	Up	0	0	AAEL009352-PA OS=Aedes aegypti GN=AAEL009352 PE=4 SV=1
CCG001783.1	1976	461	1175	10.0248217	20.78743685	1.052135305	Up	1.29E-13	3.95E-13	AAEL009394-PA OS=Aedes aegypti GN=AAEL009394 PE=4 SV=1
CCG003478.1	2249	350	1176	6.68715301	18.27965028	1.450774437	Up	3.00E-14	1.04E-13	AAEL013631-PA OS=Aedes aegypti GN=AAEL013631 PE=4 SV=1
CCG025795.1	915	113	1177	5.30664482	44.96818772	3.083032846	Up	1.42E-13	4.27E-13	AAEL011321-PA OS=Aedes aegypti GN=AAEL011321 PE=4 SV=1

CCG009379.1	11206	4137	1177	15.8634474	3.671773314	-2.111157433	Down	0	0	Ank2, isoform 1 OS=Drosophila melanogaster GN=Ank2 PE=2 SV=1
CCG006341.1	1152	151	1178	5.63231764	35.74726566	2.666032288	Up	0	0	AAEL007701-PA OS=Aedes aegypti GN=AAEL007701 PE=4 SV=1
CCG000187.1	887	209	1179	10.124774	46.46652572	2.198302075	Up	0	0	-
CCG020986.1	26840	2276	1179	3.64378226	1.535611338	-1.246623636	Down	2.78E-136	1.55E-135	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017286 PE=4 SV=1
CCG022214.1	3216	3569	1179	47.6862509	12.81586079	-1.895642979	Down	0	0	6-phosphofructokinase OS=Aedes aegypti GN=AAEL006895 PE=3 SV=1
CCG003491.1	1623	182	1180	4.81854079	25.41636882	2.399089708	Up	0	0	AAEL000437-PA OS=Aedes aegypti GN=AAEL006437 PE=4 SV=1
CCG010941.1	424	3034	1180	307.476827	97.28954385	-1.660121022	Down	1.86E-278	1.16E-277	AAEL003011-PA OS=Aedes aegypti GN=AAEL003011 PE=4 SV=1
CCG004191.1	1197	215	1181	7.71803922	34.49099822	2.159913604	Up	0	0	Ribonucleotide reductase R2 subunit OS=Aedes albopictus PE=2 SV=1
CCG009737.1	12600	396	1181	1.35047737	3.276644831	1.278749833	Up	4.78E-13	1.24E-12	AAEL010184-PA OS=Aedes aegypti GN=AAEL010184 PE=4 SV=1
CCG015570.1	3062	74	1182	1.03845864	13.49467118	3.699874158	Up	0	0	AAEL000465-PA OS=Aedes aegypti GN=AAEL008465 PE=4 SV=1
CCG004555.1	1776	312	1182	7.54873716	23.26615042	1.623925305	Up	9.28E-14	2.93E-13	AAEL000431-PA OS=Aedes aegypti GN=AAEL008431 PE=4 SV=1
CCG012046.1	2487	370	1182	6.39276309	16.61466954	1.377946063	Up	0	0	AAEL003083-PA OS=Aedes aegypti GN=AAEL003083 PE=4 SV=1
CCG005738.1	1077	373	1182	14.8818115	38.36646532	1.366295704	Up	3.38E-13	9.19E-13	AAEL000807-PA OS=Aedes aegypti GN=AAEL000807 PE=4 SV=1
CCG013727.1	1638	400	1182	10.4932197	25.22630229	1.265471334	Up	3.36E-13	9.14E-13	AAEL002867-PA OS=Aedes aegypti GN=AAEL002867 PE=3 SV=1
CCG025933.1	2163	2294	1182	45.572155	19.10341338	-1.254322152	Down	1.14E-138	6.40E-138	AAEL001616-PA OS=Aedes aegypti GN=AAEL001616 PE=3 SV=1
CCG026753.1	2178	136	1183	2.68314229	18.98789781	2.823084721	Up	1.70E-13	4.99E-13	AAEL013400-PA OS=Aedes aegypti GN=AAEL013400 PE=3 SV=1

CCG008106.1	1215	65	1184	2.29879239	34.0663372	3.889398756	Up	0	0	AAEL007075-PA (Fragment) OS=Aedes aegypti GN=AAEL007075 PE=4 SV=1
CCG022805.1	1671	399	1184	10.2602777	24.76995793	1.271521633	Up	0	0	AAEL006308-PA OS=Aedes aegypti GN=AAEL006308 PE=4 SV=1
CCG025707.1	771	156	1185	8.69426537	53.72964718	2.627582329	Up	1.42E-13	4.26E-13	AAEL013939-PA OS=Aedes aegypti GN=AAEL013939 PE=4 SV=1
CCG023565.1	2295	313	1185	5.86036032	18.05035206	1.622965701	Up	0	0	AAEL002767-PA OS=Aedes aegypti GN=AAEL002767 PE=3 SV=1
CCG001878.1	1077	420	1185	16.7569996	38.46384213	1.19873903	Up	1.84E-13	5.32E-13	AAEL002723-PA OS=Aedes aegypti GN=AAEL002723 PE=3 SV=1
CCG004351.1	752	2823	1185	161.307927	55.08717816	-1.550028866	Down	1.20E-234	7.32E-234	Mitochondrial ATP synthase F chain OS=Culex tarsalis PE=4 SV=1
CCG020696.1	1381	87	1186	2.70699994	30.02209721	3.471258003	Up	0	0	AAEL009727-PA OS=Aedes aegypti GN=AAEL009727 PE=4 SV=1
CCG012048.1	1946	111	1187	2.4509972	21.32347098	3.121001557	Up	0	0	AAEL003056-PA OS=Aedes aegypti GN=AAEL003056 PE=3 SV=1
CCG011673.1	4986	175	1187	1.50816357	8.322397619	2.464206312	Up	3.55E-13	9.57E-13	AAEL004377-PA OS=Aedes aegypti GN=AAEL004377 PE=4 SV=1
CCG025698.1	1485	271	1187	7.84161487	27.94308049	1.833268382	Up	0	0	AAEL006309-PA OS=Aedes aegypti GN=AAEL006309 PE=4 SV=1
CCG004311.2	10620	8886	1188	35.9537723	3.910586893	-3.200686101	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007110 PE=4 SV=1
CCG007435.1	1569	109	1189	2.98515046	26.49164505	3.149661879	Up	0	0	AAEL012751-PA OS=Aedes aegypti GN=AAEL012751 PE=4 SV=1
CCG007410.1	1196	3677	1189	132.106784	34.75367147	-1.926467258	Down	0	0	AAEL004042-PA OS=Aedes aegypti GN=AAEL004042 PE=2 SV=1
CCG022337.1	1611	178	1190	4.74774225	25.82268737	2.443325631	Up	1.15E-13	3.57E-13	AAEL015128-PA OS=Aedes aegypti GN=AAEL015128 PE=4 SV=1
CCG020028.1	2731	339	1190	5.33384842	15.23264349	1.513917599	Up	1.30E-13	3.98E-13	AAEL004630-PA OS=Aedes aegypti GN=AAEL004630 PE=4 SV=1
CCG018274.1	1530	103	1191	2.89273377	27.21261937	3.233770374	Up	0	0	AAEL012460-PA OS=Aedes aegypti GN=AAEL012460 PE=4 SV=1

CCG001770.1	2112	219	1191	4.4556685	19.71368733	2.145483842	Up	4.32E-13	1.14E-12	AAEL010803-PA (Fragment) OS=Aedes aegypti GN=AAEL010803 PE=3 SV=1
CCG005898.1	2132	230	1191	4.63557175	19.52875593	2.074780851	Up	3.85E-13	1.03E-12	AAEL002450-PA OS=Aedes aegypti GN=AAEL002450 PE=4 SV=1
CCG012475.1	2028	296	1191	6.27171669	20.53023059	1.710817536	Up	2.26E-13	6.47E-13	AAEL005235-PA OS=Aedes aegypti GN=AAEL005235 PE=4 SV=1
CCG021101.1	1613	306	1191	8.15172895	25.81234199	1.662883059	Up	0	0	RING finger protein 121 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006831 PE=4 SV=1
CCG003078.1	3198	433	1191	5.81797845	13.01917062	1.162047687	Up	9.43E-13	2.34E-12	AAEL009040-PA OS=Aedes aegypti GN=AAEL009040 PE=4 SV=1
CCG009392.1	2086	66	1192	1.35954098	19.97615816	3.877087605	Up	0	0	AAEL008572-PA OS=Aedes aegypti GN=AAEL008572 PE=4 SV=1
CCG005392.1	1790	223	1192	5.35321275	23.27947816	2.120581824	Up	0	0	AAEL006400-PA OS=Aedes aegypti GN=AAEL006400 PE=4 SV=1
CCG006691.1	1371	439	1193	13.7590908	30.41956542	1.144614402	Up	0	0	AAEL005492-PA OS=Aedes aegypti GN=AAEL005492 PE=4 SV=1
CCG025730.1	20349	317	1194	0.66938945	2.051215414	1.615561295	Up	2.22E-13	6.35E-13	AAEL009454-PA OS=Aedes aegypti GN=AAEL009454 PE=4 SV=1
CCG000195.1	1572	35	1195	0.95670529	26.57451701	4.79582509	Up	6.39E-14	2.11E-13	AAEL006132-PA OS=Aedes aegypti GN=AAEL006132 PE=4 SV=1
CCG022967.1	1587	215	1195	5.82135661	26.3233401	2.176915257	Up	0	0	AAEL000527-PA (Fragment) OS=Aedes aegypti GN=AAEL000527 PE=4 SV=1
CCG014156.1	3647	2856	1195	33.6500033	11.45465883	-1.554672157	Down	2.07E-238	1.26E-237	AAEL001542-PA (Fragment) OS=Aedes aegypti GN=AAEL001542 PE=4 SV=1
CCG028522.1	1506	88	1196	2.51084771	27.76234995	3.466883259	Up	2.93E-14	1.02E-13	AAEL014080-PA OS=Aedes aegypti GN=AAEL014080 PE=3 SV=1
CCG020322.1	1845	97	1196	2.25911342	22.66130028	3.326402036	Up	3.62E-14	1.25E-13	AAEL011261-PA OS=Aedes aegypti GN=AAEL011261 PE=4 SV=1
CCG010786.1	3648	99	1196	1.16611944	11.4611017	3.296958258	Up	0	0	AAEL007150-PA OS=Aedes aegypti GN=AAEL007150 PE=4 SV=1

CCG003980.1	945	119	1197	5.41100362	44.28048391	3.032702877	Up	0	0	Ribosomal protein S6 kinase, 90kD, polypeptide OS=Culex quinquefasciatus GN=CpipJ_CPIJ012176 PE=4 SV=1
CCG026970.1	1017	167	1197	7.05599379	41.1455824	2.543816348	Up	1.62E-13	4.77E-13	AAEL000737-PA OS=Aedes aegypti GN=AAEL008737 PE=4 SV=1
CCG006379.1	2075	225	1197	4.65936881	20.16629267	2.11373945	Up	0	0	AAEL002496-PA OS=Aedes aegypti GN=AAEL002496 PE=4 SV=1
CCG023834.2	1872	383	1197	8.79135062	22.3531289	1.346320059	Up	0	0	Translucan-associated protein subunit alpha OS=Culex quinquefasciatus GN=CpipJ_CPIJ018234 PE=4 SV=1
CCG024394.1	21735	2579	1197	5.09864024	1.925238431	-1.405075417	Down	2.55E-185	1.50E-184	AAEL012849-PA OS=Aedes aegypti GN=AAEL012849 PE=4 SV=1
CCG009734.1	1827	147	1199	3.45733497	22.94196708	2.730256802	Up	7.02E-14	2.29E-13	AAEL010182-PA OS=Aedes aegypti GN=AAEL010182 PE=4 SV=1
CCG007177.1	1517	224	1199	6.34490478	27.63017393	2.122574225	Up	2.04E-13	5.88E-13	mRNA capping enzyme OS=Culex quinquefasciatus GN=CpipJ_CPIJ018510 PE=4 SV=1
CCG005197.1	2139	276	1199	5.54448189	19.5955932	1.82140469	Up	0	0	AAEL011502-PA OS=Aedes aegypti GN=AAEL011502 PE=4 SV=1
CCG024240.1	3486	2395	1199	29.5216622	12.02380202	-1.295880793	Down	3.39E-152	1.93E-151	AGAP011396-PA (Fragment) OS=Anopheles gambiae GN=AGAP011396 PE=4 SV=4
CCG017753.1	1437	299	1200	8.94081465	29.19271547	1.70713022	Up	4.80E-13	1.25E-12	G-protein signaling modulator OS=Culex quinquefasciatus GN=CpipJ_CPIJ010095 PE=4 SV=1
CCG010775.1	480	4338	1200	388.338977	87.39569193	-2.151682443	Down	0	0	Conserved possible Golgi protein OS=Aedes aegypti PE=2 SV=1
CCG018527.1	3690	6901	1200	80.3615552	11.36854529	-2.821457823	Down	0	0	AAEL003104-PA OS=Aedes aegypti GN=AAEL003104 PE=4 SV=1
CCG006283.1	12053	471	1201	1.67914586	3.483356044	1.05275039	Up	0	0	AAEL000399-PA OS=Aedes aegypti GN=AAEL000399 PE=4 SV=1
CCG018360.1	1689	178	1204	4.52848594	24.91993205	2.46019945	Up	1.15E-13	3.58E-13	AAEL000716-PA OS=Aedes aegypti GN=AAEL008716 PE=4 SV=1

CCG026857.1	3664	243	1204	2.84979408	11.48738134	2.011120377	Up	1.53E-14	5.41E-14	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014388 PE=4 SV=1
CCG018727.1	2111	212	1205	4.3152931	19.95486666	2.20921018	Up	0	0	AAEL015048-PA OS=Aedes aegypti GN=AAEL015048 PE=4 SV=1
CCG027222.1	1299	247	1205	8.1705346	32.42857853	1.988763403	Up	0	0	AAEL011050-PA OS=Aedes aegypti GN=AAEL011050 PE=4 SV=1
CCG004758.1	1086	6	1206	0.23740185	38.82106979	7.353364895	Up	8.38E-06	1.47E-05	Leucine rich protein OS=Aedes albopictus PE=2 SV=1
CCG000906.1	1860	186	1207	4.29697346	22.68529036	2.400364353	Up	6.43E-13	1.64E-12	AAEL004832-PA OS=Aedes aegypti GN=AAEL004832 PE=4 SV=1
CCG000641.1	3256	312	1207	4.117493	12.95904179	1.654120946	Up	9.28E-14	2.94E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014927 PE=4 SV=1
CCG021827.1	2745	39	1208	0.61049896	15.38418883	4.655315724	Up	7.33E-14	2.38E-13	AAEL009787-PA OS=Aedes aegypti GN=AAEL009787 PE=4 SV=1
CCG003657.2	1785	454	1208	10.9289969	23.65803829	1.114169456	Up	3.16E-13	8.66E-13	AAEL011055-PA OS=Aedes aegypti GN=AAEL011055 PE=3 SV=1
CCG017249.1	2178	2611	1208	51.5123862	19.38916361	-1.409668799	Down	1.60E-188	9.45E-188	AAEL010523-PA OS=Aedes aegypti GN=AAEL010523 PE=4 SV=1
CCG023641.1	2355	119	1209	2.17129445	17.94673317	3.04709397	Up	0	0	AAEL008834-PA OS=Aedes aegypti GN=AAEL008834 PE=4 SV=1
CCG013351.1	2886	318	1209	4.73471088	14.64468351	1.629028778	Up	1.20E-13	3.69E-13	AAEL008244-PA (Fragment) OS=Aedes aegypti GN=AAEL008244 PE=4 SV=1
CCG002331.1	1420	75	1210	2.26952824	29.78839077	3.714285845	Up	0	0	AAEL000142-PA (Fragment) OS=Aedes aegypti GN=AAEL000142 PE=4 SV=1
CCG004897.1	2737	170	1210	2.66892762	15.45470036	2.5337136	Up	3.53E-13	9.53E-13	AAEL011682-PA OS=Aedes aegypti GN=AAEL011682 PE=4 SV=1
CCG027220.1	2829	238	1210	3.61498651	14.95210848	2.048286773	Up	6.86E-13	1.73E-12	AAEL011049-PA OS=Aedes aegypti GN=AAEL011049 PE=4 SV=1
CCG013174.1	716	361	1210	21.6649081	59.07753477	1.447249509	Up	1.35E-13	4.08E-13	AAEL000677-PA OS=Aedes aegypti GN=AAEL000677 PE=4 SV=1

CCG015917.1	35411	2354	1210	2.85647836	1.194530369	-1.257794069	Down	7.38E-143	4.16E-142	AAEL008347-PA OS=Aedes aegypti GN=AAEL008347 PE=4 SV=1
CCG001496.1	294	15	1211	2.1923334	143.9948067	6.037405758	Up	1.48E-08	2.92E-08	AAEL001511-PA OS=Aedes aegypti GN=AAEL001511 PE=4 SV=1
CCG025119.1	1078	219	1211	8.72947299	39.27131092	2.169509294	Up	4.32E-13	1.14E-12	AAEL005872-PA OS=Aedes aegypti GN=AAEL005872 PE=4 SV=1
CCG019267.1	2040	382	1211	8.04629344	20.75219273	1.366867525	Up	6.17E-13	1.58E-12	AAEL010060-PA OS=Aedes aegypti GN=AAEL010060 PE=4 SV=1
CCG021593.1	1116	300	1212	11.5510039	37.96544037	1.716668497	Up	1.10E-13	3.42E-13	AAEL008153-PB OS=Aedes aegypti GN=AAEL008153 PE=4 SV=1
CCG021825.3	1644	444	1212	11.6049648	25.77216025	1.151071321	Up	6.43E-13	1.64E-12	Arginyl-tRNA--protein transferase 1 OS=Aedes aegypti GN=AAEL005885 PE=3 SV=1
CCG007989.1	1190	315	1213	11.3743415	35.63394095	1.647469021	Up	4.28E-13	1.13E-12	Golgi SNAP receptor complex member 1 OS=Aedes aegypti GN=AAEL006945 PE=3 SV=1
CCG018968.1	5419	316	1213	2.50570883	7.825131892	1.642896291	Up	3.23E-13	8.83E-13	ADAM metalloprotease OS=Culex quinquefasciatus GN=CpipJ_CPIJ012681 PE=4 SV=1
CCG000066.1	3021	392	1213	5.57568221	14.03654079	1.331967195	Up	0	0	AAEL007523-PA OS=Aedes aegypti GN=AAEL007523 PE=4 SV=1
CCG024447.1	1077	14067	1213	561.239793	39.37269241	-3.833350026	Down	0	0	AAEL011112-PA OS=Aedes aegypti GN=AAEL011112 PE=3 SV=1
CCG008741.1	954	234	1214	10.5397462	44.48568973	2.07750119	Up	4.60E-13	1.20E-12	Ecdysteroid receptor OS=Aedes albopictus GN=EcR PE=2 SV=1
CCG017332.1	3984	178	1215	1.91983252	10.66122146	2.473320371	Up	1.15E-13	3.57E-13	AAEL003605-PA OS=Aedes aegypti GN=AAEL003605 PE=4 SV=1
CCG025634.1	1776	270	1215	6.53256101	23.91571299	1.872238205	Up	2.59E-13	7.26E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_15285 PE=4 SV=1
CCG019732.1	531	5888	1215	476.470428	79.98927736	-2.574508154	Down	0	0	AAEL010659-PB OS=Aedes aegypti GN=AAEL010659 PE=3 SV=1
CCG020698.1	1948	99	1216	2.18378015	21.82200439	3.320884097	Up	0	0	AAEL009725-PA OS=Aedes aegypti GN=AAEL009725 PE=4 SV=1

CCG003779.1	1722	430	1216	10.729957	24.68598406	1.202047868	Up	6.22E-13	1.59E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006642 PE=4 SV=1
CCG023810.1	4077	2332	1216	24.5782245	10.42660401	-1.237111356	Down	6.30E-138	3.53E-137	AAEL0006958-PA OS=Aedes aegypti GN=AAEL006958 PE=4 SV=1
CCG007441.1	777	88	1218	4.86658513	54.79946089	3.493180003	Up	2.93E-14	1.02E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010852 PE=4 SV=1
CCG012023.1	7671	3846	1218	21.5436839	5.550668897	-1.956531426	Down	0	0	AAEL014367-PA OS=Aedes aegypti GN=AAEL014367 PE=3 SV=1
CCG013130.1	1112	342	1219	13.2155119	38.32206779	1.5359431	Up	1.09E-13	3.41E-13	AAEL000450-PA OS=Aedes aegypti GN=AAEL006450 PE=4 SV=1
CCG015869.2	7989	418	1219	2.24825999	5.334101813	1.246436482	Up	6.74E-13	1.70E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018589 PE=4 SV=1
CCG014837.1	2008	201	1220	4.30125332	21.23959047	2.303926945	Up	0	0	AAEL011709-PA (Fragment) OS=Aedes aegypti GN=AAEL011709 PE=4 SV=1
CCG005195.1	2967	205	1220	2.96892336	14.37448522	2.275498537	Up	1.83E-13	5.31E-13	AAEL004441-PA OS=Aedes aegypti GN=AAEL004441 PE=4 SV=1
CCG010365.1	321	2580	1220	345.364222	132.8632326	-1.378176714	Down	4.95E-180	2.90E-179	AAEL000544-PA OS=Aedes aegypti GN=AAEL000544 PE=4 SV=1
CCG027711.1	1272	26	1221	0.87831219	33.55664775	5.255720971	Up	4.60E-12	1.05E-11	AAEL009654-PA OS=Aedes aegypti GN=AAEL009654 PE=4 SV=1
CCG010718.1	2911	27	1221	0.3985513	14.66302162	5.201273187	Up	2.28E-12	5.41E-12	AAEL009469-PA OS=Aedes aegypti GN=AAEL009469 PE=4 SV=1
CCG012676.1	1065	73	1221	2.94534331	40.07892576	3.76633613	Up	0	0	AAEL001402-PA OS=Aedes aegypti GN=AAEL001402 PE=4 SV=1
CCG017157.1	4275	162	1221	1.62832679	9.984574489	2.616310686	Up	0	0	AAEL005031-PA OS=Aedes aegypti GN=AAEL005031 PE=4 SV=1
CCG010559.1	636	452	1222	30.5382391	67.16826135	1.137162811	Up	1.12E-13	3.49E-13	AAEL012875-PA OS=Aedes aegypti GN=AAEL012875 PE=2 SV=1
CCG027929.1	1824	49	1223	1.15434046	23.4396779	4.343812048	Up	0	0	AAEL015128-PA OS=Aedes aegypti GN=AAEL015128 PE=4 SV=1

CCG014219.1	1668	145	1223	3.73537861	25.63187799	2.778612802	Up	0	0	Ubiquitin ligase ES OS=Culex quinquefasciatus GN=CpipJ_CPIJ009489 PE=4 SV=1
CCG014726.1	3286	166	1223	2.17071696	13.0109472	2.583482461	Up	1.83E-13	5.32E-13	AAEL009738-PA OS=Aedes aegypti GN=AAEL009738 PE=4 SV=1
CCG014393.1	1064	304	1223	12.277067	40.18230497	1.710594379	Up	0	0	Proteasome subunit alpha type OS=Aedes aegypti GN=AAEL005830 PE=3 SV=1
CCG005888.1	1317	2155	1224	70.3111451	32.48969686	-1.113771107	Down	4.29E-108	2.30E-107	26S proteasome regulatory chain 4 OS=Aedes aegypti GN=AAEL012095 PE=2 SV=1
CCG010960.2	655	31	1226	2.0336821	65.43335469	5.007860157	Up	2.40E-13	6.77E-13	Deoxyuridine 5-triphosphate nucleotidohydrolase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005616 PE=4 SV=1
CCG028044.1	1437	39	1226	1.16619321	29.8252243	4.676654249	Up	7.33E-14	2.38E-13	WD repeat protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009423 PE=4 SV=1
CCG018832.1	4024	458	1226	4.89069047	10.65080699	1.122852679	Up	1.14E-12	2.79E-12	Peroxisome biogenesis factor 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017662 PE=3 SV=1
CCG015755.1	1414	6548	1226	198.98573	30.31035879	-2.714782139	Down	0	0	AAEL006992-PA OS=Aedes aegypti GN=CYP6AG6 PE=3 SV=1
CCG018618.1	1307	156	1227	5.1287518	32.81851997	2.677830519	Up	1.42E-13	4.26E-13	AAEL011696-PA OS=Aedes aegypti GN=AAEL011696 PE=4 SV=1
CCG008136.1	1734	157	1227	3.89056997	24.73691211	2.668611989	Up	0	0	AAEL000257-PA OS=Aedes aegypti GN=AAEL000257 PE=4 SV=1
CCG007888.1	2599	345	1227	5.70394707	16.50396522	1.532780186	Up	9.59E-13	2.38E-12	-
CCG028158.1	8380	2317	1227	11.8807727	5.118592554	-1.214809591	Down	3.34E-133	1.86E-132	AAEL011164-PA (Fragment) OS=Aedes aegypti GN=AAEL011164 PE=4 SV=1
CCG016913.1	1974	222	1230	4.83246256	21.78251288	2.172339938	Up	0	0	AAEL011744-PA OS=Aedes aegypti GN=AAEL011744 PE=4 SV=1
CCG020835.1	2491	313	1230	5.39924807	17.26161398	1.676736957	Up	0	0	AAEL005180-PA OS=Aedes aegypti GN=AAEL005180 PE=4 SV=1

CCG010522.1	1667	6924	1230	178.47777	25.79404945	-2.790634205	Down	0	0	Mitochondrial NADH:ubiquinone oxidoreductase NDUFV1/51 kDa subunit OS=Aedes aegypti PE=2 SV=1
CCG003993.1	8462	121	1231	0.61443369	5.085516274	3.049065013	Up	0	0	AAEL005388-PA OS=Aedes aegypti GN=AAEL005388 PE=4 SV=1
CCG021069.1	1014	188	1231	7.96677525	42.4394859	2.413339399	Up	1.80E-13	5.25E-13	AAEL013408-PA OS=Aedes aegypti GN=AAEL013408 PE=4 SV=1
CCG020124.1	1547	279	1231	7.74955136	27.81747816	1.843806938	Up	0	0	AAEL005103-PA OS=Aedes aegypti GN=AAEL005103 PE=4 SV=1
CCG022001.1	1212	104	1232	3.68717195	35.53514603	3.268660026	Up	0	0	AAEL007084-PA OS=Aedes aegypti GN=AAEL007084 PE=4 SV=1
CCG027258.1	1194	112	1233	4.03066187	36.10013003	3.162915366	Up	0	0	AAEL001321-PA OS=Aedes aegypti GN=AAEL001321 PE=3 SV=1
CCG020505.1	2709	162	1233	2.56961868	15.91124225	2.630420285	Up	0	0	AGAP007646-PA OS=Anopheles gambiae GN=AGAP007646 PE=4 SV=2
CCG027937.1	2047	206	1233	4.3242625	21.05693955	2.283769761	Up	1.10E-13	3.43E-13	-
CCG020030.1	1323	415	1233	13.4787905	32.58016271	1.273302762	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_08793 PE=4 SV=1
CCG023520.1	1686	2440	1234	62.18633	25.58630696	-1.281225549	Down	2.59E-152	1.48E-151	AAEL005895-PA OS=Aedes aegypti GN=AAEL005895 PE=4 SV=1
CCG011210.1	897	0	1235	0.001	48.13096077	15.5546776	Up	4.44E-16	1.63E-15	AAEL013295-PA OS=Aedes aegypti GN=AAEL013295 PE=4 SV=1
CCG009601.1	3846	82	1235	0.91615139	11.22555169	3.615056526	Up	0	0	AAEL005571-PA OS=Aedes aegypti GN=AAEL005571 PE=4 SV=1
CCG006084.1	2101	129	1235	2.63831307	20.54901086	2.961381275	Up	0	0	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG011213.1	1560	209	1235	5.75684265	27.67530244	2.265249398	Up	0	0	AAEL014718-PA OS=Aedes aegypti GN=AAEL014718 PE=4 SV=1
CCG008635.1	1473	342	1235	9.97667973	29.30989261	1.554756015	Up	1.09E-13	3.41E-13	AAEL008505-PA OS=Aedes aegypti GN=AAEL008505 PE=4 SV=1
CCG019220.1	1923	391	1235	8.73695592	22.45110339	1.361583733	Up	9.02E-13	2.25E-12	AAEL014130-PA OS=Aedes aegypti GN=AAEL014130 PE=4 SV=1

CCG001764.1	1188	393	1235	14.2147354	36.34130624	1.354223028	Up	1.08E-13	3.37E-13	Suppressor of defective silencing OS=Culex quinquefasciatus GN=CpipJ_CPIJ016520 PE=4 SV-1
CCG027316.1	1566	461	1235	12.6494557	27.5692668	1.12398559	Up	1.29E-13	3.95E-13	AAEL013596-PA OS=Aedes aegypti GN=AAEL013596 PE=4 SV-1
CCG000709.1	2244	247	1236	4.7297346	19.25509362	2.025409	Up	0	0	AAEL003147-PA OS=Aedes aegypti GN=AAEL003147 PE=4 SV-1
CCG011530.1	5720	315	1237	2.36634028	7.560032931	1.67573497	Up	4.28E-13	1.13E-12	AAEL010892-PA OS=Aedes aegypti GN=AAEL010892 PE=4 SV-1
CCG013162.2	1818	217	1238	5.12895072	23.8054712	2.214557571	Up	1.47E-13	4.37E-13	AAEL004053-PA OS=Aedes aegypti GN=AAEL004053 PE=4 SV-1
CCG016709.1	2541	284	1238	4.80259922	17.03201363	1.826361684	Up	0	0	AAEL002215-PA OS=Aedes aegypti GN=AAEL002215 PE=4 SV-1
CCG001358.1	2454	257	1240	4.50009038	17.66432893	1.97281306	Up	4.14E-13	1.10E-12	AAEL009084-PA OS=Aedes aegypti GN=AAEL009084 PE=4 SV-1
CCG007394.1	2601	158	1241	2.61023378	16.67943924	2.675819856	Up	0	0	AAEL004075-PA OS=Aedes aegypti GN=AAEL004075 PE=4 SV-1
CCG002923.2	3888	227	1241	2.50877823	11.15823598	2.153052117	Up	0	0	AAEL002488-PA OS=Aedes aegypti GN=AAEL002488 PE=4 SV-1
CCG004257.1	1212	305	1241	10.8133408	35.79473719	1.726935171	Up	5.30E-13	1.37E-12	AAEL010640-PA OS=Aedes aegypti GN=AAEL010640 PE=3 SV-1
CCG018559.2	2800	471	1241	7.22812321	15.49400767	1.100017354	Up	0	0	AAEL002190-PA OS=Aedes aegypti GN=AAEL002190 PE=4 SV-1
CCG028369.2	4449	280	1242	2.70432135	9.759087379	1.851479645	Up	0	0	AAEL012168-PA OS=Aedes aegypti GN=AAEL012168 PE=4 SV-1
CCG011094.1	825	373	1243	19.4275285	52.67047034	1.438891965	Up	3.38E-13	9.18E-13	AAEL007825-PA OS=Aedes aegypti GN=AAEL007825 PE=4 SV-1
CCG028240.1	756	90	1244	5.1154446	57.52393691	3.491230878	Up	0	0	AAEL008784-PA OS=Aedes aegypti GN=AAEL008784 PE=3 SV-1
CCG012725.1	1951	101	1244	2.22447114	22.290157	3.324872491	Up	0	0	AAEL004798-PA OS=Aedes aegypti GN=AAEL004798 PE=4 SV-1
CCG006924.1	1645	339	1244	8.85516111	26.43653271	1.577942511	Up	1.30E-13	3.98E-13	AAEL008673-PA OS=Aedes aegypti GN=AAEL008673 PE=3 SV-1
CCG016095.1	933	63	1245	2.90149333	46.64850437	4.006963307	Up	0	0	AAEL006712-PA OS=Aedes aegypti GN=AAEL006712 PE=4 SV-1

CCG025009.1	2215	417	1245	8.08956178	19.64923457	1.280339657	Up	0	0	AAEL015108-FA OS=Aedes SV-1 aegypti GN=AAEL015108 PE=4
CCG014368.1	1221	3047	1245	107.230779	35.64541735	-1.588930554	Down	1.86E-262	1.15E-261	AAEL014863-FA OS=Aedes SV-1 aegypti GN=AAEL014863 PE=4
CCG015620.1	744	0	1246	0.001	58.54571621	15.83727599	Up	4.44E-16	1.65E-15	AAEL007855-FA OS=Aedes SV-1 aegypti GN=AAEL007855 PE=4
CCG019672.1	1572	34	1246	0.92937085	27.70865958	4.897938716	Up	8.82E-14	2.80E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001997 PE=4 SV-1
CCG027410.1	1662	110	1246	2.84396559	26.20819065	3.204041843	Up	0	0	AAEL002821-FA OS=Aedes SV-1 aegypti GN=AAEL002821 PE=4
CCG016916.1	240	414	1246	74.1227922	181.4917202	1.291914599	Up	8.70E-13	2.17E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007378 PE=4 SV-1
CCG020769.1	4601	4144	1246	38.7017127	9.46707517	-2.031406731	Down	0	0	AAEL005937-FA OS=Aedes SV-1 aegypti GN=AAEL005937 PE=3
CCG020439.2	3065	200	1247	2.80389785	14.22282908	2.342702764	Up	0	0	DNA ligase 4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005161 PE=3 SV-1
CCG001753.1	1742	453	1247	11.1741044	25.0246677	1.163191714	Up	9.90E-13	2.45E-12	AAEL004497-FA OS=Aedes SV-1 aegypti GN=AAEL004497 PE=4
CCG013482.2	7623	2359	1247	13.2973375	5.718610932	-1.21740075	Down	4.91E-136	2.75E-135	AAEL007465-FA OS=Aedes SV-1 aegypti GN=AAEL007465 PE=4
CCG017975.1	4875	8609	1248	75.8823477	8.949318854	-3.083914529	Down	0	0	AAEL004099-FA OS=Aedes SV-1 aegypti GN=AAEL004099 PE=4
CCG003397.1	2707	458	1249	7.27009178	16.12962235	1.149667177	Up	1.14E-12	2.79E-12	AAEL005543-FA OS=Aedes SV-1 aegypti GN=AAEL005543 PE=4
CCG008473.2	37097	2689	1249	3.11468896	1.176992417	-1.403983074	Down	6.55E-193	3.89E-192	AAEL005627-FA OS=Aedes SV-1 aegypti GN=AAEL005627 PE=4
CCG027371.1	1479	84	1250	2.44047174	29.5455348	3.597708161	Up	0	0	AAEL012708-FA OS=Aedes SV-1 aegypti GN=AAEL012708 PE=4
CCG003553.1	1293	206	1250	6.84591286	33.79570454	2.303525056	Up	1.10E-13	3.43E-13	AAEL012235-FA OS=Aedes SV-1 aegypti GN=AAEL012235 PE=2
CCG002417.1	1497	2484	1251	71.3004815	29.21363009	-1.28727018	Down	3.70E-156	2.12E-155	AAEL001965-FA OS=Aedes SV-1 aegypti GN=AAEL001965 PE=3

CCG007041.1	828	410	1252	21.277284	52.85961657	1.312851951	Up	1.25E-12	3.04E-12	AAEL014104-PA OS=Aedes aegypti GN=AAEL014104 PE=4 SV=1
CCG009676.1	1263	2574	1252	87.5725233	34.65381039	-1.337464287	Down	1.43E-171	8.29E-171	AAEL002917-PA OS=Aedes aegypti GN=AAEL002917 PE=4 SV=1
CCG023046.1	1059	149	1253	6.04578891	41.36234258	2.774315383	Up	2.33E-13	6.62E-13	AAEL004570-PA OS=Aedes aegypti GN=AAEL004570 PE=4 SV=1
CCG001911.1	2946	126	1254	1.83780942	14.88040702	3.017354913	Up	1.34E-13	4.06E-13	AAEL004364-PA OS=Aedes aegypti GN=AAEL004364 PE=4 SV=1
CCG016237.1	2100	57	1255	1.16632137	20.89173207	4.162894838	Up	0	0	AAEL004326-PA OS=Aedes aegypti GN=AAEL004328 PE=4 SV=1
CCG023053.1	1766	45	1257	1.09492529	24.88253335	4.506229042	Up	0	0	AAEL000422-PA OS=Aedes aegypti GN=AAEL000422 PE=4 SV=1
CCG022900.1	9351	477	1257	2.19191139	4.699235793	1.100236682	Up	0	0	AAEL005229-PA OS=Aedes aegypti GN=AAEL003229 PE=4 SV=1
CCG027328.1	5559	372	1258	2.87547064	7.911047343	1.4600706	Up	1.17E-13	3.60E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_13311 PE=4 SV=1
CCG023429.1	2011	125	1259	2.67091836	21.88586298	3.034591487	Up	2.35E-14	8.23E-14	AAEL000988-PA OS=Aedes aegypti GN=AAEL000988 PE=4 SV=1
CCG018220.1	1343	75	1261	2.39965011	32.82381758	3.773847074	Up	0	0	AAEL001192-PA OS=Aedes aegypti GN=AAEL001192 PE=4 SV=1
CCG027718.1	2937	155	1261	2.26772518	15.00932482	2.726541359	Up	3.42E-13	9.26E-13	Transporter OS=Aedes aegypti GN=AAEL006800 PE=3 SV=1
CCG010384.1	975	375	1261	16.526821	45.21270463	1.451918979	Up	0	0	AAEL009846-PA OS=Aedes aegypti GN=AAEL009846 PE=4 SV=1
CCG013775.1	1260	2096	1264	71.4798125	35.06925543	-1.02732905	Down	2.32E-92	1.21E-91	AAEL011081-PA OS=Aedes aegypti GN=AAEL011081 PE=4 SV=1
CCG009079.1	2259	35	1265	0.66575507	19.57601599	4.877951856	Up	6.39E-14	2.11E-13	AAEL000681-PA OS=Aedes aegypti GN=AAEL006681 PE=4 SV=1
CCG018202.1	984	113	1265	4.93453253	44.94128061	3.187055911	Up	1.42E-13	4.27E-13	AAEL000621-PA OS=Aedes aegypti GN=AAEL008621 PE=4 SV=1
CCG015568.1	4719	224	1265	2.03967378	9.371099834	2.199879951	Up	2.04E-13	5.88E-13	AAEL000462-PA OS=Aedes aegypti GN=AAEL008462 PE=4 SV=1
CCG014998.1	2371	2257	1265	40.9037077	18.65129486	-1.13295583	Down	3.03E-116	1.65E-115	AAEL002014-PA OS=Aedes aegypti GN=AAEL002014 PE=4 SV=1

CCG020504.1	888	84	1266	4.06470463	49.83916486	3.61605747	Up	0	0	Glucosamine-6-phosphate isomerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ008074 PE=3 SV=1
CCG010517.1	2754	207	1266	3.22975129	16.07014466	2.314887936	Up	6.88E-15	2.47E-14	AAEL009789-PA OS=Aedes aegypti GN=AAEL009789 PE=4 SV=1
CCG022853.1	1651	2344	1266	61.006092	26.80628613	-1.186381961	Down	8.42E-130	4.67E-129	AAEL005224-PA OS=Aedes aegypti GN=AAEL005224 PE=4 SV=1
CCG016625.1	2595	76	1267	1.25845851	17.06826076	3.7615865	Up	0	0	AAEL002195-PA (Fragment) OS=Aedes aegypti GN=AAEL002195 PE=4 SV=1
CCG026045.1	3486	167	1267	2.05850421	12.70571907	2.62580972	Up	1.62E-13	4.78E-13	AAEL003391-PA OS=Aedes aegypti GN=AAEL003391 PE=4 SV=1
CCG016807.1	4608	391	1267	3.64608642	9.612008826	1.398489216	Up	9.02E-13	2.25E-12	AAEL007449-PA OS=Aedes aegypti GN=AAEL007449 PE=4 SV=1
CCG025284.1	1455	166	1268	4.90238897	30.46535735	2.635612803	Up	1.83E-13	5.32E-13	AAEL007035-PA OS=Aedes aegypti GN=AAEL007035 PE=3 SV=1
CCG004320.1	786	480	1268	26.2410593	56.3957951	1.103761638	Up	5.41E-13	1.39E-12	AAEL006699-PA OS=Aedes aegypti GN=AAEL006699 PE=4 SV=1
CCG009256.1	3051	105	1269	1.47880109	14.54016822	3.29754404	Up	0	0	AAEL014341-PA OS=Aedes aegypti GN=AAEL014341 PE=4 SV=1
CCG006555.3	1985	341	1269	7.38170252	22.34864142	1.598161629	Up	2.98E-13	8.25E-13	AAEL000122-PA OS=Aedes aegypti GN=AAEL000122 PE=4 SV=1
CCG012545.1	1201	490	1269	17.5313655	36.93759636	1.075151619	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008696 PE=4 SV=1
CCG020476.1	2750	289	1270	4.51572847	16.14436782	1.838000303	Up	0	0	AAEL001395-PA OS=Aedes aegypti GN=AAEL001395 PE=4 SV=1
CCG010963.1	1339	387	1270	12.4191839	33.15684205	1.416736229	Up	1.31E-12	3.18E-12	AAEL000452-PA OS=Aedes aegypti GN=AAEL000452 PE=4 SV=1
CCG013643.1	600	11466	1270	821.151628	73.99501917	-3.472148578	Down	0	0	AAEL009964-PA OS=Aedes aegypti GN=AAEL009964 PE=4 SV=1
CCG001808.2	3684	55	1271	0.64151341	12.06079527	4.232701805	Up	0	0	AAEL003893-PA OS=Aedes aegypti GN=AAEL003893 PE=4 SV=1

CCG011857.2	4951	397	1271	3.44556345	8.974342512	1.381066322	Up	8.02E-13	2.01E-12	AAEL014756-PA OS=Aedes aegypti GN=AAEL014756 PE=4 SV=1
CCG019182.1	1515	20	1272	0.56725722	29.35110763	5.693268064	Up	3.69E-10	7.73E-10	Cytochrome P450 6CB1 (Fragment) OS=Aedes aegypti GN=CYP6CB1 PE=2 SV=1
CCG016776.1	2385	24	1272	0.43239985	18.64441428	5.430233658	Up	1.93E-11	4.23E-11	DNA-directed RNA polymerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ002457 PE=3 SV=1
CCG013502.1	2265	513	1272	9.73221804	19.63219782	1.012381143	Up	0	0	AAEL012514-PA OS=Aedes aegypti GN=AAEL012514 PE=4 SV=1
CCG018608.1	1052	31	1273	1.26621842	42.30217332	5.062133597	Up	2.40E-13	6.76E-13	AAEL005000-PA OS=Aedes aegypti GN=AAEL005000 PE=4 SV=1
CCG002640.1	1218	74	1273	2.61064069	36.53685249	3.806876542	Up	0	0	AAEL014441-PA OS=Aedes aegypti GN=AAEL014441 PE=4 SV=1
CCG005777.1	2013	130	1273	2.77499528	22.10724607	2.993962095	Up	0	0	AAEL007726-PA OS=Aedes aegypti GN=AAEL007726 PE=4 SV=1
CCG000340.1	1647	290	1274	7.56601277	27.04119284	1.837553676	Up	1.62E-13	4.78E-13	AAEL003670-PA OS=Aedes aegypti GN=AAEL003670 PE=4 SV=1
CCG022903.1	1698	178	1275	4.50448337	26.24958945	2.542861305	Up	1.15E-13	3.57E-13	AAEL010822-PA OS=Aedes aegypti GN=AAEL010822 PE=4 SV=1
CCG022741.1	2694	187	1277	2.98268017	16.57079415	2.473961554	Up	3.31E-13	9.03E-13	AAEL013896-PA (Fragment) OS=Aedes aegypti GN=AAEL013896 PE=4 SV=1
CCG005030.1	3405	201	1277	2.5365394	13.11063713	2.369804322	Up	0	0	AAEL013108-PA OS=Aedes aegypti GN=AAEL013108 PE=4 SV=1
CCG011949.1	3081	362	1277	5.04869975	14.48936041	1.521010126	Up	1.06E-12	2.62E-12	AAEL008413-PA (Fragment) OS=Aedes aegypti GN=AAEL008413 PE=4 SV=1
CCG007984.3	5709	2535	1277	19.0800976	7.819533971	-1.286914018	Down	2.96E-159	1.70E-158	AAEL004972-PA OS=Aedes aegypti GN=AAEL004972 PE=4 SV=1
CCG028207.1	1456	176	1278	5.19414374	30.6845314	2.562553706	Up	0	0	AAEL011317-PA OS=Aedes aegypti GN=AAEL011317 PE=3 SV=1
CCG009510.1	1650	184	1278	4.79177647	27.07677437	2.498423369	Up	3.33E-13	9.08E-13	AGAP006107-PC OS=Anopheles gambiae GN=AGAP006107 PE=4 SV=2

CCG007638.1	1002	209	1278	8.96274904	44.58750271	2.314626193	Up	0	0	GI21485 OS=Drosophila mojavensis GN=GI21485 PE=4 SV=1
CCG018049.1	1014	263	1278	11.1450101	44.05983996	1.983066335	Up	1.40E-14	4.97E-14	-
CCG010363.1	4995	2119	1278	18.2288023	8.944279823	-1.027182548	Down	2.48E-93	1.30E-92	AAEL001582-PA (Fragment) OS=Aedes aegypti GN=AAEL001582 PE=3 SV=1
CCG004929.1	1210	2226	1278	79.0501068	36.92287414	-1.098252553	Down	4.11E-109	2.20E-108	AAEL013739-PA OS=Aedes aegypti GN=AAEL013739 PE=4 SV=1
CCG006605.1	438	7047	1278	691.341826	102.0015473	-2.760808174	Down	0	0	AAEL001164-PA OS=Aedes aegypti GN=AAEL001164 PE=4 SV=1
CCG020230.3	2876	224	1279	3.34673872	15.54646592	2.215758831	Up	2.04E-13	5.88E-13	ES ubiquitin ligase protein CPT OS=Culex quinquefasciatus GN=CpipJ_CPIJ002840 PE=4 SV=1
CCG026027.1	1412	487	1279	14.820298	31.66546458	1.095335791	Up	0	0	AAEL012586-PA OS=Aedes aegypti GN=AAEL012586 PE=4 SV=1
CCG025267.1	2253	343	1280	6.54177495	19.86089404	1.602176533	Up	0	0	AAEL008944-PA OS=Aedes aegypti GN=AAEL008944 PE=4 SV=1
CCG012226.3	6504	395	1280	2.60963179	6.879857667	1.398532456	Up	0	0	AGAP007503-PA OS=Anopheles gambiae GN=AGAP007503 PE=4 SV=1
CCG013756.1	3931	209	1281	2.2845776	11.39189838	2.318008832	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009801 PE=4 SV=1
CCG011664.1	1287	79	1282	2.63761386	34.82246373	3.722713002	Up	0	0	AAEL011535-PA OS=Aedes aegypti GN=AAEL011535 PE=4 SV=1
CCG022815.1	1511	268	1282	7.62136921	29.660166	1.96040456	Up	0	0	AAEL004384-PA OS=Aedes aegypti GN=AAEL004384 PE=4 SV=1
CCG025161.1	1389	353	1282	10.9203141	32.26530657	1.562969377	Up	0	0	AAEL015607-PA OS=Aedes aegypti GN=AAEL015607 PE=4 SV=1
CCG011941.3	2544	164	1283	2.77006151	17.63029446	2.670066654	Up	1.14E-13	3.54E-13	AAEL006488-PA (Fragment) OS=Aedes aegypti GN=AAEL006488 PE=4 SV=1
CCG027099.1	3187	2756	1283	37.1586409	14.0732567	-1.400741514	Down	5.65E-197	3.36E-196	-

CCG025042.1	1327	390	1284	12.6286334	33.82549162	1.421412377	Up	6.59E-13	1.68E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001780 PE=4 SV=1
CCG024571.1	3260	500	1284	6.59045009	13.76884275	1.062958406	Up	2.39E-13	6.76E-13	AAEL002788-PA OS=Aedes aegypti GN=AAEL002788 PE=4 SV=1
CCG004887.1	3079	84	1285	1.17228246	14.58960236	3.637548425	Up	0	0	Chromosome 10 SCAF15019, whole genome shotgun sequence. (Fragment) OS=Tetraodon nigroviridis GN=GSTENG00032246001 PE=4 SV=1
CCG024094.1	1068	240	1286	9.65612014	42.09395499	2.124097536	Up	0	0	AAEL013816-PA OS=Aedes aegypti GN=AAEL013816 PE=4 SV=1
CCG025399.1	2661	282	1286	4.5537261	16.89452985	1.891436779	Up	5.45E-13	1.40E-12	AAEL005827-PA (Fragment) OS=Aedes aegypti GN=AAEL005827 PE=4 SV=1
CCG027517.1	1953	304	1286	6.68858132	23.01912132	1.783060618	Up	0	0	Anoctamin OS=Aedes aegypti GN=AAEL010243 PE=3 SV=1
CCG017362.1	3059	149	1287	2.09300113	14.70784642	2.812941022	Up	2.33E-13	6.63E-13	AAEL008183-PA OS=Aedes aegypti GN=AAEL008183 PE=4 SV=1
CCG024208.1	2879	308	1287	4.59697057	15.62740612	1.765323001	Up	0	0	AAEL010078-PA OS=Aedes aegypti GN=AAEL010078 PE=4 SV=1
CCG012816.1	4878	2685	1287	23.6518527	9.223309185	-1.358596831	Down	3.13E-183	1.84E-182	AGAP007865-PA (Fragment) OS=Anopheles gambiae GN=AGAP007865 PE=4 SV=4
CCG017636.2	1848	5565	1287	129.397496	24.34594275	-2.41005643	Down	0	0	AAEL014609-PA (Fragment) OS=Aedes aegypti GN=CYP9J26 PE=3 SV=1
CCG017679.1	1857	49	1288	1.13382714	24.24677463	4.418520238	Up	0	0	AAEL007797-PA OS=Aedes aegypti GN=AAEL007797 PE=4 SV=1
CCG025419.1	4716	86	1288	0.78358719	9.547553113	3.606965327	Up	0	0	AAEL009460-PA OS=Aedes aegypti GN=AAEL009460 PE=4 SV=1
CCG016871.2	1108	117	1288	4.53741782	40.63741921	3.162865362	Up	9.44E-14	2.97E-13	AAEL005312-PE OS=Aedes aegypti PE=2 SV=1

CCG013390.1	1946	135	1289	2.98094253	23.15581642	2.957534155	Up	1.90E-13	5.48E-13	AAEL002483-PA OS=Aedes aegypti GN=AAEL002483 PE=4 SV=1
CCG007489.1	3306	221	1289	2.87244747	13.63013272	2.246447193	Up	0	0	AAEL001719-PA OS=Aedes aegypti GN=AAEL001719 PE=4 SV=1
CCG010751.1	1593	65	1290	1.75331623	28.30896236	4.013100741	Up	0	0	AAEL000676-PA OS=Aedes aegypti GN=AAEL006876 PE=4 SV=1
CCG006716.1	1743	179	1290	4.41284136	25.87273496	2.551652777	Up	0	0	AAEL004874-PA OS=Aedes aegypti GN=AAEL004874 PE=4 SV=1
CCG002033.1	1671	384	1290	9.87455302	26.98753862	1.450506053	Up	1.03E-12	2.55E-12	AAEL0006367-PA OS=Aedes aegypti GN=AAEL006367 PE=4 SV=1
CCG001069.1	1857	514	1290	11.8936153	24.2844249	1.029844005	Up	1.57E-13	4.65E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006377 PE=4 SV=1
CCG019307.1	975	0	1291	0.001	46.28834391	15.49836133	Up	4.44E-16	1.63E-15	AAEL013715-PA OS=Aedes aegypti GN=AAEL013715 PE=3 SV=1
CCG026849.1	2827	130	1291	1.9759694	15.96432095	3.014218676	Up	0	0	AAEL002215-PA OS=Aedes aegypti GN=AAEL002215 PE=4 SV=1
CCG020784.1	2166	243	1292	4.8207043	20.85230544	2.112891055	Up	1.53E-14	5.41E-14	AAEL009815-PA OS=Aedes aegypti GN=AAEL009815 PE=4 SV=1
CCG020471.1	2242	144	1293	2.75987591	20.16104008	2.868894762	Up	7.53E-14	2.44E-13	Fanconi anemia group J protein OS=Danaus plexippus GN=KGM_13519 PE=4 SV=1
CCG004926.1	1123	222	1293	8.49446223	40.2502688	2.244403897	Up	0	0	AAEL007568-PA OS=Aedes aegypti GN=AAEL007568 PE=4 SV=1
CCG012191.1	840	259	1293	13.2490015	53.81077603	2.022011476	Up	0	0	AAEL009244-PA OS=Aedes aegypti GN=AAEL009244 PE=3 SV=1
CCG002221.1	1728	127	1294	3.15807656	26.17824661	3.051250419	Up	1.57E-13	4.66E-13	AAEL010349-PA OS=Aedes aegypti GN=AAEL010349 PE=4 SV=1
CCG022486.1	1641	146	1294	3.82302331	27.5661244	2.850110547	Up	9.33E-15	3.34E-14	AAEL012185-PA OS=Aedes aegypti GN=AAEL012185 PE=2 SV=1
CCG004980.1	1724	236	1294	5.88216785	26.238985	2.157292056	Up	4.60E-13	1.20E-12	AAEL003811-PA OS=Aedes aegypti GN=AAEL003811 PE=4 SV=1
CCG005800.1	918	314	1294	14.6977088	49.2766995	1.745314357	Up	4.97E-14	1.68E-13	AAEL011848-PA OS=Aedes aegypti GN=AAEL011848 PE=4 SV=1

CCG023983.1	2146	86	1295	1.72199309	21.09551185	3.614784832	Up	0	0	AGAP004519-PA OS=Anopheles gambiae GN=AGAP004519 PE=4 SV=4
CCG020606.1	1968	105	1295	2.29259255	23.00354086	3.326804069	Up	0	0	AAEL002151-PA OS=Aedes aegypti GN=AAEL002151 PE=4 SV=1
CCG024827.1	1319	440	1295	14.334104	34.32218986	1.259689873	Up	0	0	AAEL011399-PA OS=Aedes aegypti GN=AAEL011399 PE=4 SV=1
CCG015943.1	3120	3406	1295	46.908627	14.50992578	-1.692813133	Down	0	0	AAEL006085-PA OS=Aedes aegypti GN=AAEL006085 PE=3 SV=1
CCG009970.1	2565	224	1297	3.75252263	17.67675827	2.235921046	Up	2.04E-13	5.88E-13	Zinc finger protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ012784 PE=3 SV=1
CCG006714.1	543	411	1298	32.5240533	83.56508886	1.361393288	Up	0	0	Signal recognition particle 9 kDa protein OS=Aedes aegypti GN=AAEL012049 PE=3 SV=1
CCG025759.1	1029	51	1300	2.1296953	44.16497551	4.37418377	Up	0	0	AAEL011085-PA OS=Aedes aegypti GN=AAEL011085 PE=4 SV=1
CCG006535.1	2232	111	1300	2.13693573	20.3610035	3.252193245	Up	0	0	AAEL007869-PA OS=Aedes aegypti GN=AAEL007869 PE=4 SV=1
CCG027413.1	1605	183	1300	4.89935292	28.31511514	2.530909273	Up	3.05E-13	8.41E-13	AAEL000706-PA OS=Aedes aegypti GN=AAEL000706 PE=3 SV=1
CCG003009.1	1130	173	1301	6.57855229	40.24842308	2.613090223	Up	3.60E-13	9.69E-13	AAEL000782-PA OS=Aedes aegypti GN=AAEL000782 PE=4 SV=1
CCG000956.1	456	194	1301	18.2809836	99.73841684	2.447805608	Up	5.97E-13	1.53E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013272 PE=4 SV=1
CCG014027.1	1712	238	1301	5.97359628	26.56584	2.152900687	Up	6.86E-13	1.73E-12	Transcription factor Spo OS=Culex quinquefasciatus GN=CpipJ_CPIJ005037 PE=4 SV=1
CCG021450.1	2679	241	1302	3.86551177	16.98980081	2.135937601	Up	2.37E-13	6.72E-13	Pita OS=Culex quinquefasciatus GN=CpipJ_CPIJ010913 PE=4 SV=1
CCG009333.1	2069	384	1302	7.97504983	21.99887692	1.463864436	Up	1.03E-12	2.55E-12	Asparagine synthetase OS=Aedes aegypti GN=AAEL015631 PE=4 SV=1

CCG011511.1	2250	205	1303	3.91502026	20.2447265	2.370454473	Up	1.83E-13	5.31E-13	AAEL007048-PA OS=Aedes aegypti GN=AAEL007048 PE=4 SV=1
CCG020279.1	1586	493	1304	13.3569225	28.74249238	1.105597522	Up	6.97E-14	2.27E-13	AAEL008480-PA OS=Aedes aegypti GN=AAEL008480 PE=4 SV=1
CCG017940.1	6822	2796	1304	17.6111665	6.682144959	-1.398107287	Down	3.11E-199	1.85E-198	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014431 PE=4 SV=1
CCG013061.1	588	2	1305	0.14615556	77.58597141	9.052147295	Up	9.36E-05	0.00015325	AAEL008182-PA OS=Aedes aegypti GN=AAEL008182 PE=4 SV=1
CCG003013.1	2745	22	1305	0.34438403	16.61950863	5.592715677	Up	8.38E-11	1.80E-10	AAEL009404-PA OS=Aedes aegypti GN=AAEL009404 PE=4 SV=1
CCG002741.1	2053	100	1305	2.09302166	22.22140827	3.408291106	Up	0	0	AAEL003542-PA OS=Aedes aegypti GN=AAEL003542 PE=4 SV=1
CCG022129.1	1560	105	1305	2.89219368	29.24394307	3.337901778	Up	0	0	Alkaline phosphatase OS=Aedes aegypti GN=AAEL003298 PE=3 SV=1
CCG019369.1	2098	221	1305	4.52636385	21.74478131	2.264244736	Up	0	0	Chitin synthase OS=Aedes aegypti PE=2 SV=2
CCG014967.1	1824	263	1305	6.19574573	25.0112671	2.013228306	Up	1.40E-14	4.97E-14	AAEL006544-PA OS=Aedes aegypti GN=AAEL006544 PE=4 SV=1
CCG006067.1	1188	489	1305	17.6870372	38.40113736	1.11845664	Up	1.91E-13	5.51E-13	AAEL007043-PA OS=Aedes aegypti GN=AAEL007043 PE=4 SV=1
CCG009701.1	975	511	1305	22.5205481	46.79030891	1.054967814	Up	0	0	AAEL010731-PA OS=Aedes aegypti GN=AAEL010731 PE=4 SV=1
CCG022205.1	1431	108	1306	3.24299884	31.90461877	3.298364883	Up	0	0	AAEL003604-PA OS=Aedes aegypti GN=AAEL003604 PE=4 SV=1
CCG022417.1	857	434	1306	21.7606357	53.27363998	1.291701153	Up	0	0	-
CCG019493.1	912	244	1307	11.4962886	50.09919708	2.123619292	Up	2.78E-13	7.73E-13	AAEL004285-PA OS=Aedes aegypti GN=AAEL004285 PE=4 SV=1
CCG022189.1	848	254	1307	12.8706516	53.88026856	2.065671943	Up	2.35E-13	6.67E-13	-
CCG008941.1	642	515	1307	34.4694912	71.16895287	1.045928008	Up	9.44E-13	2.34E-12	AAEL013128-PA OS=Aedes aegypti GN=AAEL013128 PE=4 SV=1
CCG021807.1	750	83	1308	4.7553173	60.96723469	3.680420598	Up	1.32E-13	4.02E-13	Lysozyme OS=Aedes albopictus PE=2 SV=1

CCG013295.1	1871	444	1308	10.1969867	24.43903047	1.261044163	Up	6.43E-13	1.64E-12	AAEL014720-PA OS=Aedes aegypti GN=AAEL014720 PE=4 SV=1
CCG023758.1	1104	133	1309	5.17660752	41.44962346	3.00128015	Up	9.90E-14	3.11E-13	AAEL003970-PA (Fragment) OS=Aedes aegypti GN=AAEL003970 PE=4 SV=1
CCG003775.1	757	425	1309	24.1243556	60.449649	1.325243555	Up	1.58E-12	3.80E-12	AAEL013761-PA OS=Aedes aegypti GN=AAEL013761 PE=3 SV=1
CCG027651.2	1854	334	1310	7.74104173	24.70083202	1.673960008	Up	3.15E-13	8.65E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000129 PE=4 SV=1
CCG025520.1	13752	7615	1311	23.7939594	3.332628043	-2.835863147	Down	0	0	Dynein heavy chain, cytosolic, putative OS=Pediculus humanus subsp. corporis GN=Phum_PHUM561990 PE=4 SV=1
CCG026317.1	9864	337	1312	1.46804547	4.649762685	1.663260427	Up	0	0	AAEL003771-PA OS=Aedes aegypti GN=AAEL003771 PE=4 SV=1
CCG006461.1	359	428	1312	51.2285415	127.758382	1.318398222	Up	1.52E-13	4.51E-13	AAEL005951-PA OS=Aedes aegypti GN=AAEL005951 PE=4 SV=1
CCG011696.1	1584	15	1314	0.40691037	28.9994796	6.155172169	Up	1.48E-08	2.92E-08	Importin subunit alpha OS=Aedes aegypti GN=AAEL012960 PE=3 SV=1
CCG010050.1	1914	155	1314	3.4797852	23.99956932	2.785938359	Up	3.42E-13	9.25E-13	AAEL005265-PA OS=Aedes aegypti GN=AAEL005265 PE=4 SV=1
CCG001411.1	2685	481	1314	7.69774389	17.10807288	1.15216968	Up	0	0	AAEL003768-PA OS=Aedes aegypti GN=AAEL003768 PE=4 SV=1
CCG023849.1	1488	132	1315	3.8118313	30.89390723	3.018766169	Up	1.46E-13	4.36E-13	AAEL015615-PA OS=Aedes aegypti GN=AAEL015615 PE=4 SV=1
CCG002711.1	2109	90	1316	1.83370133	21.81369949	3.572403881	Up	0	0	AAEL007578-PA OS=Aedes aegypti GN=AAEL007578 PE=4 SV=1
CCG007874.3	10660	2386	1316	9.61780364	4.315674693	-1.15612135	Down	8.51E-127	4.71E-126	Receptor protein-tyrosine phosphatase 10d OS=Culex quinquefasciatus GN=CpipJ_CPIJ006052 PE=4 SV=1

CCG012586.1	1865	219	1317	5.04577581	24.68635416	2.290565774	Up	4.32E-13	1.14E-12	AAEL003690-PA OS=Aedes aegypti GN=AAEL003690 PE=4 SV=1
CCG017299.1	2710	308	1317	4.88364511	16.98894853	1.798566293	Up	0	0	inhibitor of kappa kinase epsilon OS=Culex quinquefasciatus GN=CpipJ_CPIJ006852 PE=4 SV=1
CCG000473.1	774	2301	1317	127.743358	59.48326939	-1.102692434	Down	1.87E-113	1.01E-112	Larval chymotrypsin-like protein (Precursor) OS=Aedes aegypti PE=2 SV=1
CCG016335.2	3828	4886	1317	54.8459047	12.02718143	-2.189085314	Down	0	0	AAEL003104-PA OS=Aedes aegypti GN=AAEL003104 PE=4 SV=1
CCG024446.1	1908	130	1318	2.92770728	24.14832746	3.044080046	Up	0	0	AAEL011124-PA OS=Aedes aegypti GN=AAEL011124 PE=4 SV=1
CCG008714.1	816	219	1320	11.5323185	56.5501536	2.293848358	Up	4.32E-13	1.14E-12	AAEL002705-PA OS=Aedes aegypti GN=AAEL002705 PE=4 SV=1
CCG005019.1	2485	379	1320	6.55353296	18.56938645	1.50258138	Up	1.40E-13	4.20E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000685 PE=4 SV=1
CCG008043.1	4455	136	1321	1.31175845	10.36585491	2.982265114	Up	1.70E-13	4.98E-13	AAEL001361-PA (Fragment) OS=Aedes aegypti GN=AAEL001361 PE=4 SV=1
CCG026822.1	2949	467	1321	6.80463413	15.65950614	1.202449215	Up	5.22E-14	1.75E-13	AAEL000730-PA OS=Aedes aegypti GN=AAEL000730 PE=4 SV=1
CCG020323.1	3201	526	1321	7.06094358	14.42670528	1.030808966	Up	1.08E-13	3.37E-13	AAEL011267-PA OS=Aedes aegypti GN=AAEL011267 PE=4 SV=1
CCG008705.1	2868	2775	1321	41.5763646	16.10177253	-1.368544101	Down	2.00E-191	1.18E-190	AAEL003194-PA OS=Aedes aegypti GN=AAEL003194 PE=4 SV=1
CCG023623.1	1809	403	1322	9.57258322	25.54717628	1.416183637	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_15504 PE=4 SV=1
CCG003743.1	2325	15	1323	0.27722409	19.89238717	6.165019954	Up	1.48E-08	2.93E-08	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002513 PE=4 SV=1
CCG018789.1	2630	112	1323	1.82988984	17.58547535	3.264555628	Up	0	0	AAEL013688-PA OS=Aedes aegypti GN=AAEL013688 PE=4 SV=1

CCG025867.1	4446	201	1325	1.94262633	10.41828986	2.423038157	Up	0	0	AAEL007270-PA OS=Aedes aegypti GN=AAEL007270 PE=4 SV=1
CCG028311.1	4848	266	1325	2.35766283	9.554397014	2.018807413	Up	4.09E-14	1.40E-13	AAEL003391-PA OS=Aedes aegypti GN=AAEL003391 PE=4 SV=1
CCG027696.1	482	6	1327	0.53489296	96.24405244	7.491303358	Up	8.38E-06	1.47E-05	Predicted protein OS=Nematostella vectensis GN=v1g229551 PE=3 SV=1
CCG011239.1	975	81	1328	3.56979334	47.6149657	3.737502632	Up	4.46E-14	1.51E-13	AAEL012551-PA OS=Aedes aegypti GN=AAEL012551 PE=4 SV=1
CCG020644.1	1005	339	1328	14.4942687	46.19362344	1.672211172	Up	1.30E-13	3.98E-13	AAEL011667-PA OS=Aedes aegypti GN=AAEL011667 PE=4 SV=1
CCG019086.1	2001	207	1329	4.44514496	23.21816583	2.384951636	Up	6.88E-15	2.47E-14	AAEL005495-PA (Fragment) OS=Aedes aegypti GN=AAEL005495 PE=4 SV=1
CCG001246.1	1742	435	1329	10.7301002	26.67023526	1.313567002	Up	1.05E-12	2.58E-12	AAEL000460-PA OS=Aedes aegypti GN=AAEL000460 PE=4 SV=1
CCG002024.1	1617	96	1331	2.55107886	28.7751802	3.495645559	Up	0	0	AAEL009880-PA OS=Aedes aegypti GN=AAEL009880 PE=4 SV=1
CCG011009.1	2838	2218	1331	33.5824071	16.39516081	-1.03443559	Down	1.01E-98	5.32E-98	-
CCG024742.1	2338	256	1332	4.70498377	19.9163493	2.081691571	Up	0	0	AAEL004950-PA OS=Aedes aegypti GN=AAEL004950 PE=4 SV=1
CCG014912.1	3640	53	1333	0.62565822	12.80202828	4.354853814	Up	0	0	AAEL006331-PA OS=Aedes aegypti GN=AAEL006331 PE=4 SV=1
CCG010192.1	963	324	1333	14.457107	48.38980575	1.742924266	Up	0	0	AAEL001211-PA OS=Aedes aegypti GN=AAEL001211 PE=4 SV=1
CCG014231.1	12311	385	1333	1.34378587	3.785182596	1.494059633	Up	7.93E-13	1.99E-12	AAEL003899-PA OS=Aedes aegypti GN=AAEL003899 PE=4 SV=1
CCG022381.1	1899	2434	1333	55.0754787	24.53890623	-1.166339184	Down	3.99E-131	2.22E-130	AAEL001770-PA OS=Aedes aegypti GN=AAEL001770 PE=4 SV=1
CCG024246.1	984	196	1334	8.55901218	47.39262319	2.469146311	Up	7.93E-14	2.55E-13	AAEL010411-PA OS=Aedes aegypti GN=AAEL010411 PE=4 SV=1
CCG010142.1	1523	373	1334	10.5237761	30.62005333	1.540824335	Up	3.38E-13	9.19E-13	AAEL009448-PA OS=Aedes aegypti GN=AAEL009448 PE=4 SV=1
CCG021818.1	1773	124	1336	3.00521551	26.34193896	3.131821186	Up	1.61E-13	4.76E-13	AAEL008890-PA OS=Aedes aegypti GN=AAEL008890 PE=3 SV=1

CCG022947.1	2605	203	1336	3.34850523	17.9286978	2.420681579	Up	9.28E-14	2.93E-13	AAEL011212-PA OS=Aedes aegypti GN=AAEL011212 PE=4 SV=1
CCG001365.1	2053	12991	1336	271.904443	22.74927315	-3.579207372	Down	0	0	AAEL004294-PA OS=Aedes aegypti GN=AAEL004294 PE=3 SV=1
CCG004799.1	822	72	1337	3.76377237	56.86036015	3.917171952	Up	0	0	AAEL001955-PA OS=Aedes aegypti GN=AAEL001955 PE=4 SV=1
CCG024551.1	1686	85	1338	2.16632707	27.7426894	3.678784668	Up	0	0	AAEL003339-PA OS=Aedes aegypti GN=AAEL003339 PE=4 SV=1
CCG001268.1	2273	164	1338	3.10032401	20.57816732	2.7306236	Up	1.14E-13	3.54E-13	Canal debranching enzyme OS=Culex quinquefasciatus GN=CpipJ_CPIJ004547 PE=4 SV=1
CCG003229.1	11395	477	1338	1.79873308	4.104798098	1.190330148	Up	0	0	AAEL007980-PA OS=Aedes aegypti GN=AAEL007980 PE=4 SV=1
CCG016773.1	1650	412	1339	10.7294125	28.36917127	1.402752922	Up	3.41E-13	9.25E-13	AAEL001886-PA OS=Aedes aegypti GN=AAEL001886 PE=4 SV=1
CCG006246.1	1050	10078	1339	412.427605	44.58012628	-3.209668292	Down	0	0	AAEL003430-PA OS=Aedes aegypti GN=AAEL003430 PE=4 SV=1
CCG018636.1	3634	255	1340	3.01521253	12.89050382	2.095977052	Up	0	0	AAEL004526-PA OS=Aedes aegypti GN=AAEL004526 PE=4 SV=1
CCG009558.1	1137	3717	1340	140.473618	41.19972812	-1.769592484	Down	0	0	AAEL006194-PA OS=Aedes aegypti GN=AAEL006194 PE=4 SV=1
CCG004352.1	1212	4750	1340	168.404488	38.65024	-2.123381309	Down	0	0	AAEL006194-PA OS=Aedes aegypti GN=AAEL006194 PE=4 SV=1
CCG003310.1	1008	325	1341	13.8543291	46.50699321	1.747110818	Up	1.19E-13	3.66E-13	AAEL011106-PA OS=Aedes aegypti GN=AAEL011106 PE=4 SV=1
CCG007653.4	2725	498	1341	7.85281755	17.20332079	1.131404794	Up	3.42E-13	9.26E-13	AAEL006031-PA OS=Aedes aegypti GN=AAEL006031 PE=4 SV=1
CCG005016.1	1443	55	1342	1.63779307	32.51143966	4.311122446	Up	0	0	AAEL014037-PA OS=Aedes aegypti GN=AAEL014037 PE=4 SV=1
CCG013246.1	1712	158	1342	3.96566476	27.40304172	2.788701412	Up	0	0	Ryndremine 5-monooxygenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ017147 PE=4 SV=1
CCG024831.1	318	210	1342	28.3762398	147.5283252	2.378236642	Up	0	0	AAEL005884-PA OS=Aedes aegypti GN=AAEL005884 PE=4 SV=1

CCG023673.3	2040	277	1342	5.83461593	22.99706246	1.978739994	Up	0	0	AAEL002080-PA OS=Aedes aegypti GN=AAEL002080 PE=4 SV=1
CCG019717.1	1904	355	1342	8.01168896	24.63970978	1.620806946	Up	2.28E-13	6.52E-13	AAEL003292-PA (Fragment) OS=Aedes aegypti GN=AAEL003292 PE=4 SV=1
CCG003239.1	2043	399	1342	8.39203334	22.96329292	1.452237224	Up	0	0	AAEL010860-PA (Fragment) OS=Aedes aegypti GN=AAEL010860 PE=4 SV=1
CCG011500.1	2886	123	1344	1.83135044	16.27994594	3.152116121	Up	0	0	AAEL007863-PA OS=Aedes aegypti GN=AAEL007863 PE=4 SV=1
CCG028409.1	394	320	1345	34.8992768	119.3372646	1.773775566	Up	0	0	AAEL004284-PA OS=Aedes aegypti GN=AAEL004284 PE=4 SV=1
CCG007914.1	1572	208	1348	5.68556285	29.97694471	2.398478267	Up	3.93E-13	1.05E-12	AAEL003514-PA OS=Aedes aegypti GN=AAEL003514 PE=4 SV=1
CCG021007.1	1637	315	1348	8.2684584	28.78665674	1.799709967	Up	4.28E-13	1.13E-12	AAEL004464-PA OS=Aedes aegypti GN=AAEL004464 PE=4 SV=1
CCG003714.1	805	483	1350	25.7818408	58.62568154	1.185177517	Up	0	0	AAEL002435-PA OS=Aedes aegypti GN=AAEL002435 PE=4 SV=1
CCG014954.1	2324	3971	1350	73.4220379	20.30708849	-1.85422975	Down	0	0	AGAP007051-PA OS=Anopheles gambiae GN=AGAP007051 PE=4 SV=4
CCG024096.1	1485	68	1351	1.96763768	31.80379254	4.014662322	Up	0	0	AAEL013820-PA OS=Aedes aegypti GN=AAEL013820 PE=4 SV=1
CCG010912.1	3316	350	1351	4.53540625	14.24265136	1.650914051	Up	3.00E-14	1.04E-13	AAEL008382-PA OS=Aedes aegypti GN=AAEL008382 PE=4 SV=1
CCG027000.1	2169	333	1352	6.5970132	21.79049802	1.723814273	Up	4.41E-13	1.16E-12	Transcriptional repressor protein YY1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014689 PE=4 SV=1
CCG022351.1	1965	382	1352	8.35340388	24.05271766	1.525763812	Up	6.17E-13	1.58E-12	AAEL001506-PA OS=Aedes aegypti GN=AAEL001506 PE=4 SV=1
CCG016275.1	1803	415	1352	9.89042699	26.21386034	1.406225114	Up	0	0	AAEL010587-PA OS=Aedes aegypti GN=AAEL010587 PE=4 SV=1
CCG014838.1	1764	101	1353	2.46028526	26.8132361	3.446047845	Up	0	0	AAEL011707-PA OS=Aedes aegypti GN=AAEL011707 PE=4 SV=1

CCG001345.1	5892	145	1353	1.05746971	8.027587996	2.924350238	Up	0	0	AAEL008618-PA OS=Aedes aegypti GN=AAEL008618 PE=4 SV=1
CCG024235.1	879	203	1353	9.92361334	53.80949769	2.438923411	Up	9.28E-14	2.93E-13	AAEL010642-PA OS=Aedes aegypti GN=AAEL010642 PE=4 SV=1
CCG021034.1	5904	248	1353	1.80496175	8.01127176	2.150063017	Up	5.86E-14	1.95E-13	AAEL008557-PA (Fragment) OS=Aedes aegypti GN=AAEL008557 PE=4 SV=1
CCG012987.1	1871	31	1354	0.71195178	25.29850708	5.151128917	Up	2.40E-13	6.77E-13	AAEL001666-PA OS=Aedes aegypti GN=AAEL001666 PE=4 SV=1
CCG009896.1	1842	40	1354	0.93311042	25.69680062	4.783397133	Up	6.86E-14	2.24E-13	AAEL004698-PA OS=Aedes aegypti GN=AAEL004698 PE=4 SV=1
CCG014519.1	2109	91	1354	1.85407579	22.44357835	3.597530587	Up	0	0	AAEL007578-PA OS=Aedes aegypti GN=AAEL007578 PE=4 SV=1
CCG008609.1	1587	454	1354	12.2925391	29.82577615	1.27877674	Up	3.16E-13	8.66E-13	AAEL010321-PA OS=Aedes aegypti GN=AAEL010321 PE=3 SV=1
CCG020032.1	2130	493	1354	9.94557707	22.22230364	1.159881391	Up	6.97E-14	2.28E-13	AAEL004633-PA OS=Aedes aegypti GN=AAEL004633 PE=4 SV=1
CCG024079.1	2025	89	1355	1.88854636	23.39183458	3.630656909	Up	0	0	AAEL006549-PA OS=Aedes aegypti GN=AAEL006549 PE=4 SV=1
CCG017175.1	1389	97	1355	3.0007662	34.10256661	3.506477498	Up	3.62E-14	1.25E-13	AAEL011309-PA OS=Aedes aegypti GN=AAEL011309 PE=3 SV=1
CCG027741.1	780	219	1357	12.0645793	60.81843792	2.33373115	Up	4.32E-13	1.14E-12	Putative salivary secreted peptide of the 13 kDa family OS=Aedes triseriatus PE=2 SV=1
CCG018182.1	1424	314	1357	9.47506788	33.31347021	1.81389746	Up	4.97E-14	1.68E-13	AAEL002581-PA OS=Aedes aegypti GN=AAEL002581 PE=4 SV=1
CCG003545.1	3810	411	1357	4.63531783	12.45101879	1.425523626	Up	0	0	AAEL002061-PA OS=Aedes aegypti GN=AAEL002061 PE=3 SV=1
CCG017655.1	1821	420	1357	9.91064719	26.05073124	1.394272692	Up	1.84E-13	5.32E-13	AAEL000064-PA OS=Aedes aegypti GN=DCE PE=2 SV=1
CCG007433.1	1203	457	1357	16.3234985	39.43340115	1.272467854	Up	1.13E-12	2.77E-12	AAEL004108-PA OS=Aedes aegypti GN=AAEL004108 PE=4 SV=1
CCG001121.2	2220	516	1357	9.98755994	21.36864035	1.097290954	Up	9.75E-14	3.07E-13	AAEL005734-PA (Fragment) OS=Aedes aegypti GN=AAEL005734 PE=4 SV=1

CCG017935.1	1185	262	1358	9.50048141	40.06189017	2.076157967	Up	0	0	AAEL003645-PA OS=Aedes aegypti GN=AAEL003645 PE=3 SV=1
CCG001635.1	3090	498	1358	6.92521936	15.36354041	1.149579036	Up	3.42E-13	9.27E-13	AAEL001205-PA OS=Aedes aegypti GN=Aael_AAEL001205 PE=4 SV=1
CCG006050.2	4747	502	1358	4.54409243	10.00070357	1.138037414	Up	6.14E-13	1.57E-12	AAEL001244-PA OS=Aedes aegypti GN=AAEL001244 PE=4 SV=1
CCG002109.1	2406	538	1358	9.60836127	19.7312302	1.038118606	Up	1.07E-13	3.33E-13	AAEL003379-PA OS=Aedes aegypti GN=AAEL003379 PE=4 SV=1
CCG002419.1	15520	4728	1358	13.0902645	3.058849218	-2.097433352	Down	0	0	AAEL001963-PA OS=Aedes aegypti GN=AAEL001963 PE=4 SV=1
CCG008436.1	1446	420	1359	12.4808358	32.85497796	1.396397427	Up	1.84E-13	5.32E-13	AAEL002562-PA OS=Aedes aegypti GN=AAEL002562 PE=4 SV=1
CCG014944.1	939	470	1359	21.5077479	50.5945667	1.234125998	Up	1.05E-12	2.59E-12	AAEL003680-PA OS=Aedes aegypti GN=AAEL003680 PE=4 SV=1
CCG020662.1	1331	2391	1359	77.1905601	35.69368755	-1.11275547	Down	1.52E-119	8.33E-119	AAEL004042-PA OS=Aedes aegypti GN=AAEL004042 PE=2 SV=1
CCG025660.1	2068	252	1361	5.23615722	23.00687364	2.135484632	Up	5.42E-14	1.81E-13	AAEL001331-PA OS=Aedes aegypti GN=AAEL001331 PE=2 SV=1
CCG022002.1	2235	414	1362	7.95949446	21.30343309	1.420337234	Up	8.70E-13	2.17E-12	AAEL007100-PA OS=Aedes aegypti GN=AAEL007100 PE=4 SV=1
CCG024119.1	802	5776	1362	309.467814	59.36804609	-2.382030835	Down	0	0	AAEL013744-PA OS=Aedes aegypti GN=AAEL013744 PE=2 SV=1
CCG000603.1	5355	32	1365	0.25677526	8.910933295	5.11699844	Up	1.77E-13	5.16E-13	AAEL005656-PA OS=Aedes aegypti GN=AAEL005656 PE=4 SV=1
CCG018725.1	861	507	1365	25.3027357	55.4216583	1.131156503	Up	8.62E-13	2.15E-12	AAEL004983-PA OS=Aedes aegypti GN=AAEL004983 PE=4 SV=1
CCG008281.1	1158	4	1366	0.14842741	41.23748365	8.118054972	Up	2.95E-05	5.04E-05	AAEL000828-PA OS=Aedes aegypti GN=AAEL000828 PE=3 SV=1
CCG022465.1	1152	77	1366	2.872109	41.45226221	3.851268431	Up	0	0	Cen growth-regulating nuclear protein OS=Culex quinquefasciatus GN=Cqnl103225 PE=4 SV=1
CCG019480.1	1854	280	1366	6.48949606	25.75674545	1.988771955	Up	0	0	AAEL000589-PA OS=Aedes aegypti GN=AAEL000589 PE=4 SV=1
CCG026086.1	876	7794	1366	382.312913	54.51256401	-2.810093244	Down	0	0	AAEL003235-PA OS=Aedes aegypti GN=AAEL003235 PE=4 SV=1

CCG002509.1	2301	62	1367	1.15781119	20.76834609	4.164914421	Up	0	0	AGAP011312-PA (Fragment) OS=Anopheles gambiae GN=AGAP011312 PE=4 SV=4
CCG021608.1	1062	193	1367	7.80900073	44.99808319	2.526653694	Up	6.08E-14	2.02E-13	AAEL0008392-PA OS=Aedes aegypti GN=AAEL008392 PE=4 SV=1
CCG025894.1	2546	164	1370	2.7678855	18.81101303	2.764721377	Up	1.14E-13	3.54E-13	Zinc finger protein 169 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004616 PE=4 SV=1
CCG027415.1	588	2721	1370	198.844639	81.45040677	-1.28764786	Down	6.08E-171	3.53E-170	AAEL000713-PA OS=Aedes aegypti GN=AAEL000713 PE=4 SV=1
CCG006549.1	2433	71	1371	1.25394622	19.69905362	3.97357894	Up	0	0	AAEL000999-PA OS=Aedes aegypti GN=AAEL000999 PE=3 SV=1
CCG006647.1	1572	249	1373	6.80627476	30.53289695	2.165427182	Up	0	0	AAEL000736-PA OS=Aedes aegypti GN=AAEL005736 PE=4 SV=1
CCG009894.1	2090	230	1374	4.72872678	22.98213985	2.280989442	Up	3.85E-13	1.03E-12	AAEL000698-PA OS=Aedes aegypti GN=AAEL006988 PE=4 SV=1
CCG018420.1	1542	501	1374	13.9609838	31.14959292	1.157812699	Up	0	0	AAEL000860-PA OS=Aedes aegypti GN=AAEL000860 PE=4 SV=1
CCG010114.1	1512	10514	1374	298.798803	31.7676404	-3.233544527	Down	0	0	AAEL005617-PA OS=Aedes aegypti GN=UGP PE=2 SV=1
CCG016021.1	1970	2551	1376	55.6425345	24.41755779	-1.188269225	Down	1.92E-141	1.08E-140	AAEL002163-PA OS=Aedes aegypti GN=AAEL002163 PE=4 SV=1
CCG004750.1	1304	111	1377	3.6576998	36.91529687	3.335210182	Up	0	0	AAEL012176-PA OS=Aedes aegypti GN=AAEL012176 PE=4 SV=1
CCG022924.1	2195	195	1377	3.81735683	21.93054538	2.522295734	Up	2.26E-13	6.47E-13	AAEL002563-PA OS=Aedes aegypti GN=AAEL002563 PE=4 SV=1
CCG015556.1	4242	274	1377	2.77550855	11.34784232	2.031593965	Up	0	0	AAEL000545-PA OS=Aedes aegypti GN=AAEL005450 PE=4 SV=1
CCG025395.1	1890	329	1377	7.47991677	25.46960165	1.767682274	Up	0	0	AAEL000659-PA OS=Aedes aegypti GN=AAEL006659 PE=4 SV=1
CCG007700.1	674	351	1377	22.3774137	71.42069305	1.674298828	Up	4.44E-15	1.60E-14	AAEL000382-PA OS=Aedes aegypti GN=AAEL005382 PE=4 SV=1
CCG007765.1	2223	216	1378	4.17519689	21.67004291	2.375785874	Up	1.38E-13	4.17E-13	AAEL000944-PA OS=Aedes aegypti GN=AAEL009644 PE=4 SV=1

CCG004515.1	2860	87	1379	1.3071213	16.85575653	3.688776449	Up	0	0	AAEL004502-PA OS=Aedes aegypti GN=AAEL004562 PE=4 SV=1
CCG013887.1	1446	209	1380	6.21070161	33.36267078	2.425406623	Up	0	0	AAEL002101-PA OS=Aedes aegypti GN=AAEL002101 PE=4 SV=1
CCG016222.1	560	23	1381	1.76482839	86.20960754	5.610248852	Up	4.01E-11	8.68E-11	Deoxyuridine 5'-triphosphate nucleotidohydrolase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005616 PE=4 SV=1
CCG002102.1	629	222	1381	15.1657887	76.75259177	2.339394942	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_16009 PE=4 SV=1
CCG005367.1	621	162	1382	11.209496	77.79764654	2.795005101	Up	0	0	AAEL012993-PA OS=Aedes aegypti GN=AAEL012993 PE=4 SV=1
CCG003037.2	2430	315	1382	5.57015078	19.88162078	1.835647086	Up	4.28E-13	1.13E-12	AAEL013503-PA OS=Aedes aegypti GN=AAEL013503 PE=4 SV=1
CCG026666.1	1368	106	1384	3.32952622	35.36714551	3.409020977	Up	0	0	AAEL010402-PA OS=Aedes aegypti GN=AAEL010402 PE=3 SV=1
CCG004685.1	3468	94	1385	1.16469292	13.96113418	3.583394613	Up	0	0	AAEL009662-PA OS=Aedes aegypti GN=AAEL009662 PE=4 SV=1
CCG025667.2	3840	116	1386	1.29804407	12.61775302	3.281043751	Up	1.16E-13	3.58E-13	AAEL000984-PA (Fragment) OS=Aedes aegypti GN=AAEL000984 PE=4 SV=1
CCG005354.1	1196	278	1387	9.98794835	40.5410785	2.021124203	Up	2.11E-14	7.38E-14	S107p: RNA polymerase II holoenzyme component OS=Culex quinquefasciatus GN=CpipJ_CPIJ013518 PE=4 SV=1
CCG003402.1	2215	74	1388	1.43555773	21.90613461	3.931651691	Up	0	0	AAEL013237-PA OS=Aedes aegypti GN=AAEL013237 PE=4 SV=1
CCG007925.1	3327	89	1388	1.14947592	14.58433669	3.665371625	Up	0	0	Raptorin OS=Culex quinquefasciatus GN=CpipJ_CPIJ004643 PE=4 SV=1
CCG025206.1	1758	120	1388	2.93308769	27.60073274	3.234214461	Up	0	0	AAEL011816-PA OS=Aedes aegypti GN=AAEL011816 PE=4 SV=1
CCG003587.1	6493	2346	1388	15.5254886	7.472984469	-1.054882242	Down	1.14E-107	6.10E-107	AAEL006222-PA OS=Aedes aegypti GN=AAEL006222 PE=4 SV=1

CCG008041.1	2445	205	1390	3.60277938	19.87403056	2.463702272	Up	1.83E-13	5.30E-13	AAEL001361-PA (Fragment) OS=Aedes aegypti GN=AAEL001361 PE=4 SV=1
CCG026572.1	1462	391	1390	11.4919058	33.23666533	1.532157574	Up	9.02E-13	2.25E-12	AAEL009307-PA (Fragment) OS=Aedes aegypti GN=AAEL009307 PE=4 SV=1
CCG009046.1	1274	108	1391	3.64264626	38.16873076	3.389332406	Up	0	0	AAEL008049-PA OS=Aedes aegypti GN=AAEL008049 PE=4 SV=1
CCG023877.1	1050	537	1391	21.97595	46.31139332	1.07544163	Up	3.67E-13	9.84E-13	AAEL000090-PA OS=Aedes aegypti GN=AAEL000090 PE=4 SV=1
CCG019112.1	1860	7425	1391	171.532408	26.14352849	-2.713955402	Down	0	0	Malic enzyme OS=Aedes aegypti GN=AAEL001091 PE=3 SV=1
CCG015402.1	1623	10139	1393	268.435083	30.00423878	-3.161335001	Down	0	0	AAEL004294-PA OS=Aedes aegypti GN=AAEL004294 PE=3 SV=1
CCG020983.1	1926	245	1394	5.46603582	25.30209648	2.210690111	Up	3.40E-13	9.24E-13	AAEL002646-PA OS=Aedes aegypti GN=AAEL002646 PE=4 SV=1
CCG023952.1	1086	269	1394	10.6435162	44.87277884	2.075865687	Up	0	0	AAEL012416-PA OS=Aedes aegypti GN=AAEL012416 PE=4 SV=1
CCG001013.1	345	357	1394	44.4643341	141.2517038	1.667547786	Up	3.04E-13	8.39E-13	40S ribosomal protein S21 OS=Aedes aegypti GN=AAEL000529 PE=3 SV=1
CCG005810.1	1701	378	1395	9.54882991	28.66948624	1.586120186	Up	1.08E-12	2.65E-12	AAEL001014-PA OS=Aedes aegypti GN=AAEL001014 PE=4 SV=1
CCG027247.1	327	11953	1395	1570.69492	149.133933	-3.396722525	Down	0	0	AAEL004457-PA OS=Aedes aegypti GN=AAEL004457 PE=3 SV=1
CCG026230.1	1305	99	1396	3.25977297	37.39598036	3.52003981	Up	0	0	AAEL012100-PA OS=Aedes aegypti GN=AAEL012100 PE=4 SV=1
CCG024172.1	2572	114	1396	1.90456833	18.97424354	3.316506416	Up	8.48E-14	2.70E-13	AAEL009285-PA OS=Aedes aegypti GN=AAEL009285 PE=3 SV=1
CCG010254.1	1721	201	1396	5.01854541	28.3566266	2.498344739	Up	0	0	AGAP013497-PA OS=Anopheles gambiae GN=AgaP_AGAP013497 PE=4 SV=1
CCG023939.1	1473	376	1396	10.9685134	33.13085837	1.594807578	Up	3.71E-14	1.28E-13	AAEL003935-PA OS=Aedes aegypti GN=AAEL003935 PE=4 SV=1

CCG012769.2	5304	3264	1396	26.4429136	9.200934083	-1.523028912	Down	4.86E-264	3.01E-263	AAEL010384-PA OS=Aedes aegypti GN=AAEL010384 PE=4 SV=1
CCG016189.1	2103	191	1397	3.90262449	23.22240259	2.573000681	Up	6.68E-14	2.19E-13	AAEL011535-PA OS=Aedes aegypti GN=AAEL011535 PE=4 SV=1
CCG019812.1	2199	278	1397	5.43228114	22.20860057	2.031488437	Up	2.11E-14	7.39E-14	AAEL005020-PA OS=Aedes aegypti GN=AAEL005020 PE=4 SV=1
CCG008851.2	7118	564	1397	3.40473874	6.861016107	1.010878157	Up	9.30E-13	2.31E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017078 PE=4 SV=1
CCG021081.1	1819	306	1398	7.22855349	26.86732871	1.894074007	Up	0	0	AAEL001638-PA OS=Aedes aegypti GN=AAEL001638 PE=4 SV=1
CCG005601.1	2163	440	1398	8.74095387	22.59439248	1.370102136	Up	0	0	AAEL011636-PA OS=Aedes aegypti GN=AAEL011636 PE=4 SV=1
CCG012914.1	426	10617	1399	1070.91472	114.8042939	-3.221595096	Down	0	0	AAEL005292-PA OS=Aedes aegypti GN=AAEL005292 PE=4 SV=1
CCG006252.1	942	48	1401	2.18954062	51.99208679	4.569591944	Up	5.35E-14	1.80E-13	AAEL010593-PA OS=Aedes aegypti GN=AAEL010593 PE=4 SV=1
CCG003973.1	2960	216	1401	3.13562928	16.54613032	2.399666942	Up	1.38E-13	4.17E-13	AAEL010459-PA OS=Aedes aegypti GN=AAEL010459 PE=4 SV=1
CCG020710.1	2960	368	1402	5.34218322	16.55794055	1.632021882	Up	3.69E-13	9.90E-13	AAEL007197-PA OS=Aedes aegypti GN=AAEL007197 PE=4 SV=1
CCG009854.1	1661	391	1402	10.115091	29.50722699	1.544559041	Up	9.02E-13	2.25E-12	AAEL012460-PA OS=Aedes aegypti GN=AAEL012460 PE=4 SV=1
CCG023937.1	2026	109	1403	2.31179717	24.20852039	3.388428173	Up	0	0	AAEL014814-PA OS=Aedes aegypti GN=AAEL014814 PE=4 SV=1
CCG011599.1	1443	351	1404	10.4520976	34.01345848	1.702313204	Up	4.44E-15	1.60E-14	AAEL002101-PA OS=Aedes aegypti GN=AAEL002101 PE=4 SV=1
CCG009436.1	1176	276	1405	10.0847336	41.76562829	2.050143162	Up	0	0	AAEL004839-PA OS=Aedes aegypti GN=AAEL004839 PE=3 SV=1
CCG024360.1	1523	296	1405	8.35130758	32.24975631	1.949214253	Up	2.26E-13	6.47E-13	AAEL013261-PA OS=Aedes aegypti GN=AAEL013261 PE=4 SV=1
CCG009090.1	6237	3040	1405	20.9440425	7.87500062	-1.411187989	Down	1.62E-219	9.76E-219	AAEL010061-PA OS=Aedes aegypti GN=AAEL010061 PE=4 SV=1

CCG005936.1	4296	372	1406	3.72084294	11.44118649	1.620535272	Up	1.17E-13	3.60E-13	AAEL009490-PA (Fragment) OS=Aedes aegypti GN=AAEL009490 PE=3 SV=1
CCG001925.1	3000	469	1406	6.71760184	16.38377905	1.28624997	Up	0	0	AAEL001205-PA OS=Aedes aegypti GN=Aael_AAEL001205 PE=4 SV=1
CCG019484.1	1896	99	1407	2.24367285	25.94213893	3.531363197	Up	0	0	AAEL009789-PA OS=Aedes aegypti GN=AAEL009789 PE=4 SV=1
CCG004230.1	1227	455	1407	15.9341722	40.08663033	1.330997082	Up	0	0	AAEL009533-PA OS=Aedes aegypti GN=AAEL009533 PE=4 SV=1
CCG024493.1	2007	441	1409	9.44178025	24.54220826	1.378134255	Up	0	0	SKeritropin OS=Culex quinquefasciatus GN=CpipJ_CPIJ009263 PE=4 SV=1
CCG002684.1	4775	246	1410	2.21372874	10.32275817	2.221278146	Up	2.51E-13	7.05E-13	AAEL011960-PA OS=Aedes aegypti GN=AAEL011960 PE=4 SV=1
CCG018305.1	2865	521	1410	7.81404249	17.20459695	1.138653089	Up	0	0	AAEL002114-PA OS=Aedes aegypti GN=AAEL002114 PE=4 SV=1
CCG019255.1	3777	6060	1410	68.9427037	13.05034955	-2.401309427	Down	0	0	AAEL003129-PA OS=Aedes aegypti GN=AAEL003129 PE=4 SV=1
CCG018641.1	1901	345	1411	7.79829481	25.94746372	1.734362925	Up	9.59E-13	2.38E-12	AAEL011750-PA OS=Aedes aegypti GN=AAEL011750 PE=3 SV=1
CCG003443.1	1087	77	1412	3.04385425	45.41038344	3.899051036	Up	0	0	AAEL012855-PA OS=Aedes aegypti GN=AAEL012855 PE=4 SV=1
CCG008978.1	1162	400	1412	14.791647	42.47942066	1.521981387	Up	3.36E-13	9.14E-13	AAEL005528-PA OS=Aedes aegypti GN=AAEL005528 PE=4 SV=1
CCG016513.1	2186	476	1412	9.35663023	22.58055206	1.271019814	Up	4.62E-13	1.21E-12	AAEL013887-PA OS=Aedes aegypti GN=AAEL013887 PE=4 SV=1
CCG023389.1	1980	200	1413	4.34037723	24.94749751	2.523002764	Up	0	0	AAEL005634-PA OS=Aedes aegypti GN=AAEL005634 PE=4 SV=1
CCG002515.1	15474	431	1414	1.19684345	3.194455432	1.41633555	Up	0	0	AAEL005477-PA OS=Aedes aegypti GN=AAEL005477 PE=4 SV=1
CCG022773.2	1305	5618	1415	184.983884	37.90495144	-2.286941368	Down	0	0	Glutamine synthetase OS=Aedes aegypti GN=AAEL013458 PE=3 SV=1
CCG003985.1	1518	6984	1415	197.694747	32.58627249	-2.600938302	Down	0	0	AAEL005269-PA OS=Aedes aegypti GN=AAEL005269 PE=3 SV=1

CCG009723.1	3357	58	1416	0.74240233	14.74558234	4.311937759	Up	0	0	AAEL007071-FA OS=Aedes aegypti GN=AAEL007071 PE=4 SV=1
CCG014273.1	540	19808	1416	1576.19352	91.6683702	-4.103876845	Down	0	0	AAEL012207-FA OS=Aedes aegypti GN=AAEL012207 PE=4 SV=1
CCG016236.1	945	10542	1417	479.351262	52.41891872	-3.19292373	Down	0	0	AAEL004338-FA OS=Aedes aegypti GN=AAEL004338 PE=4 SV=1
CCG008976.1	2913	357	1418	5.26611578	17.01710829	1.692174757	Up	3.04E-13	8.39E-13	AAEL005469-FA OS=Aedes aegypti GN=AAEL005469 PE=4 SV=1
CCG024001.2	2713	344	1419	5.44842931	18.28448018	1.746707323	Up	4.69E-13	1.22E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007750 PE=4 SV=1
CCG019688.1	3884	370	1419	4.09340932	12.77183181	1.641590617	Up	0	0	AAEL005571-FA OS=Aedes aegypti GN=AAEL005571 PE=4 SV=1
CCG010407.1	3268	400	1419	5.25945344	15.17925176	1.529115888	Up	3.36E-13	9.14E-13	AAEL006026-FA OS=Aedes aegypti GN=AAEL006026 PE=4 SV=1
CCG023168.1	3366	451	1419	5.75738274	14.73731276	1.355988455	Up	1.51E-12	3.65E-12	AAEL009677-FA OS=Aedes aegypti GN=AAEL009677 PE=4 SV=1
CCG001650.3	33355	6975	1419	8.98557634	1.487207158	-2.595005424	Down	0	0	AAEL007874-FA OS=Aedes aegypti GN=AAEL007874 PE=4 SV=1
CCG011728.1	1355	469	1420	14.8729192	36.63524208	1.300544306	Up	0	0	-
CCG002126.1	1921	524	1420	11.721052	25.84109996	1.140565417	Up	0	0	AAEL014698-FA OS=Aedes aegypti GN=AAEL014698 PE=4 SV=1
CCG019419.1	1080	79	1421	3.14315651	45.99602898	3.871223295	Up	0	0	PKP36 pre-miRNA processing factor 38 domain containing B OS=Culex quinquefasciatus GN=CpipJ_CPIJ004500 PE=4 SV=1
CCG022358.1	621	82	1421	5.67394241	79.99309387	3.817452038	Up	0	0	AAEL001495-FA OS=Aedes aegypti GN=AAEL001495 PE=4 SV=1
CCG022557.1	1559	109	1421	3.00429831	31.86383021	3.406819718	Up	0	0	AAEL003984-FA OS=Aedes aegypti GN=AAEL003984 PE=4 SV=1
CCG006948.1	726	189	1421	11.1863359	68.42384476	2.612761619	Up	2.34E-13	6.64E-13	AAEL006748-FA OS=Aedes aegypti GN=AAEL006748 PE=4 SV=1
CCG015210.1	689	513	1421	31.9934308	72.09827474	1.172189027	Up	0	0	AAEL008692-FA OS=Aedes aegypti GN=AAEL008692 PE=4 SV=1

CCG014407.1	4774	3635	1421	32.7178436	10.40546948	-1.652735606	Down	0	0	AAEL005458-PA OS=Aedes aegypti GN=AAEL005458 PE=3 SV=1
CCG021273.1	37980	4898	1421	5.54148921	1.307943952	-2.082973016	Down	0	0	AAEL011520-PA OS=Aedes aegypti GN=AAEL011520 PE=4 SV=1
CCG023517.1	705	4906	1421	299.020593	70.46200183	-2.085327475	Down	0	0	Hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG003056.1	1701	502	1422	12.6812503	29.22437952	1.204475399	Up	6.14E-13	1.57E-12	AAEL000448-PA OS=Aedes aegypti GN=AAEL000448 PE=4 SV=1
CCG009951.1	8403	268	1423	1.37044971	5.919984273	2.11094396	Up	0	0	AAEL000065-PA OS=Aedes aegypti GN=AAEL000065 PE=4 SV=1
CCG025974.1	1044	329	1423	13.5412286	47.64906882	1.815089377	Up	0	0	AAEL013609-PA (Fragment) OS=Aedes aegypti GN=AAEL013609 PE=4 SV=1
CCG008020.1	1245	7120	1423	245.738563	39.95632759	-2.620628376	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_05645 PE=4 SV=1
CCG002702.1	4015	11386	1423	121.856388	12.38994467	-3.297940234	Down	0	0	AAEL009510-PA OS=Aedes aegypti GN=AAEL009510 PE=2 SV=1
CCG003435.1	1126	5	1424	0.19080699	44.21011201	7.85611854	Up	1.59E-05	2.76E-05	AAEL013721-PA OS=Aedes aegypti GN=AAEL013721 PE=4 SV=1
CCG028046.1	6406	213	1425	1.42874703	7.776388448	2.444349788	Up	3.57E-13	9.60E-13	AAEL004064-PA OS=Aedes aegypti GN=AAEL004064 PE=4 SV=1
CCG004912.1	2783	129	1426	1.99176995	17.91250545	3.168844215	Up	0	0	AAEL005152-PA OS=Aedes aegypti GN=AAEL005152 PE=4 SV=1
CCG008431.1	11109	2564	1426	9.9175803	4.487397847	-1.144109076	Down	7.28E-134	4.06E-133	AAEL008773-PA OS=Aedes aegypti GN=AAEL008773 PE=4 SV=1
CCG010176.1	945	321	1427	14.5960686	52.78884757	1.854653336	Up	0	0	AAEL006460-PA OS=Aedes aegypti GN=AAEL006460 PE=4 SV=1
CCG028088.1	1342	146	1428	4.67479974	37.19852402	2.992268909	Up	9.33E-15	3.34E-14	AAEL005904-PA OS=Aedes aegypti GN=AAEL005904 PE=4 SV=1
CCG015823.1	3426	146	1429	1.83116791	14.58125438	2.993278846	Up	9.33E-15	3.34E-14	AAEL010905-PA OS=Aedes aegypti GN=AAEL010905 PE=4 SV=1
CCG017184.1	2491	246	1429	4.24349848	20.05434665	2.2405889	Up	2.51E-13	7.05E-13	AAEL013597-PA OS=Aedes aegypti GN=AAEL013597 PE=4 SV=1

CCG013020.1	1875	184	1430	4.21676329	26.66151242	2.660550679	Up	3.33E-13	9.08E-13	AAEL010648-PA OS=Aedes aegypti GN=AAEL010648 PE=4 SV=1
CCG005696.1	3540	219	1430	2.65829714	14.12156378	2.409325576	Up	4.32E-13	1.14E-12	AAEL010568-PA OS=Aedes aegypti GN=AAEL010568 PE=3 SV=1
CCG012557.1	1257	379	1430	12.9558707	39.7695591	1.618058597	Up	1.40E-13	4.20E-13	mRNA cap guanine-N7 methyltransferase OS=Aedes aegypti GN=AAEL011698 PE=3 SV=1
CCG003329.1	2571	258	1431	4.31201538	19.45752394	2.173893905	Up	1.69E-13	4.97E-13	AAEL002260-PA (Fragment) OS=Aedes aegypti GN=AAEL002260 PE=4 SV=1
CCG017627.1	1928	305	1431	6.79759806	25.94672928	1.932455728	Up	5.30E-13	1.37E-12	Cation-transporting ATPase OS=Culex quinquefasciatus GN=CpipJ_CPIJ006431 PE=4 SV=1
CCG028458.1	4953	427	1431	3.70443704	10.0999988	1.447028901	Up	9.31E-13	2.31E-12	AAEL010443-PA OS=Aedes aegypti GN=AAEL010443 PE=4 SV=1
CCG019327.1	2640	288	1433	4.68760741	18.97545857	2.017211097	Up	0	0	AAEL007141-PA OS=Aedes aegypti GN=AAEL007141 PE=4 SV=1
CCG004956.1	1469	87	1434	2.54483792	34.12537025	3.745199017	Up	0	0	AAEL007850-PA OS=Aedes aegypti GN=AAEL007850 PE=4 SV=1
CCG016218.1	1726	151	1434	3.75922939	29.0441303	2.949737773	Up	0	0	AAEL010118-PA OS=Aedes aegypti GN=AAEL010118 PE=4 SV=1
CCG013475.1	1385	194	1435	6.01886535	36.22030842	2.589235383	Up	5.97E-13	1.53E-12	SV=1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017451 PE=4 SV=1
CCG025061.1	1419	288	1436	8.72113007	35.37708629	2.020228236	Up	0	0	AAEL004422-PA OS=Aedes aegypti GN=AAEL004422 PE=4 SV=1
CCG011329.1	2973	290	1438	4.19146419	16.90884696	2.012252074	Up	1.62E-13	4.78E-13	AAEL005848-PA OS=Aedes aegypti GN=AAEL005848 PE=4 SV=1
CCG020970.1	2022	313	1438	6.65159591	24.86152423	1.902142317	Up	0	0	AGAP001943-PA OS=Anopheles gambiae GN=AgaP_AGAP001943 PE=4 SV=5
CCG023367.1	891	513	1438	24.7401502	56.41975533	1.189346149	Up	0	0	AAEL006964-PA OS=Aedes aegypti GN=AAEL006964 PE=4 SV=1
CCG023639.1	759	78	1439	4.41586205	66.27794503	3.907761862	Up	0	0	AAEL008843-PA OS=Aedes aegypti GN=AAEL008843 PE=4 SV=1

CCG016045.1	1639	348	1439	9.1235312	30.69247119	1.750220585	Up	0	0	AAEL011361-PA OS=Aedes aegypti GN=AAEL011361 PE=4 SV=1
CCG011860.1	1458	228	1440	6.71954698	34.52669311	2.361276286	Up	1.05E-13	3.28E-13	AAEL008600-PA OS=Aedes aegypti GN=AAEL008600 PE=3 SV=1
CCG021949.1	6795	73	1441	0.46163217	7.41352124	4.005343265	Up	0	0	DNA polymerase OS=Aedes aegypti GN=AAEL002800 PE=3 SV=1
CCG013089.1	2000	98	1441	2.105517	25.18743841	3.58045798	Up	0	0	AAEL004317-PA OS=Aedes aegypti GN=AAEL004317 PE=4 SV=1
CCG010849.1	3633	273	1441	3.2289396	13.8659171	2.102410683	Up	1.53E-14	5.42E-14	AAEL013280-PA OS=Aedes aegypti GN=AAEL013280 PE=4 SV=1
CCG012553.1	2520	102	1442	1.73925116	20.00390282	3.523743311	Up	0	0	AAEL011693-PA OS=Aedes aegypti GN=AAEL011693 PE=3 SV=1
CCG019573.1	954	575	1442	25.8989491	52.84049801	1.028750507	Up	1.93E-13	5.56E-13	AAEL001761-PA OS=Aedes aegypti GN=AAEL001761 PE=4 SV=1
CCG002083.1	1867	537	1443	12.3592649	27.0191716	1.12839051	Up	3.67E-13	9.84E-13	AAEL001339-PA OS=Aedes aegypti GN=AAEL001339 PE=4 SV=1
CCG028389.1	10083	15	1445	0.06392403	5.009888916	6.292276386	Up	1.48E-08	2.92E-08	AAEL009992-PA OS=Aedes aegypti GN=AAEL009992 PE=4 SV=1
CCG013465.1	3297	381	1445	4.96556533	15.32141642	1.625519794	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011525 PE=4 SV=1
CCG000070.1	2584	426	1446	7.08401972	19.5625651	1.465455421	Up	4.37E-13	1.15E-12	AAEL006236-PA OS=Aedes aegypti GN=AAEL006236 PE=4 SV=1
CCG005981.4	1269	520	1446	17.6077715	39.83425391	1.177797228	Up	1.17E-12	2.86E-12	AAEL005405-PA OS=Aedes aegypti GN=AAEL005405 PE=4 SV=1
CCG020368.1	1416	267	1448	8.10234403	35.74829433	2.141463159	Up	5.21E-13	1.35E-12	AAEL010610-PA OS=Aedes aegypti GN=AAEL010610 PE=4 SV=1
CCG017768.1	4612	160	1450	1.49071065	10.9907852	2.882222294	Up	4.24E-14	1.44E-13	AAEL006211-PA (Fragment) OS=Aedes aegypti GN=AAEL006211 PE=4 SV=1
CCG016268.2	1562	280	1450	7.70264129	32.45166538	2.074867372	Up	0	0	AAEL007136-PA OS=Aedes aegypti GN=AAEL007136 PE=4 SV=1
CCG001751.1	2006	274	1451	5.8692459	25.28637069	2.107112925	Up	0	0	AAEL002012-PA OS=Aedes aegypti GN=AAEL002012 PE=4 SV=1

CCG019260.1	2682	232	1452	3.71699419	18.92595745	2.348157947	Up	2.91E-13	8.05E-13	AAEL009640-PA OS=Aedes aegypti GN=AAEL009640 PE=4 SV=1
CCG010973.1	2031	236	1452	4.99303662	24.99232785	2.323495892	Up	4.60E-13	1.20E-12	Condensin complex subunit 2 OS=Aedes aegypti GN=AAEL005962 PE=3 SV=1
CCG011003.1	2927	9941	1452	145.938549	17.3417895	-3.073036328	Down	0	0	AGAP001213-PA OS=Anopheles gambiae GN=AgaP_AGAP001213 PE=4 SV=5
CCG019316.1	1831	413	1453	9.69224489	27.7413305	1.51713422	Up	5.88E-13	1.51E-12	AAEL006663-PA OS=Aedes aegypti GN=AAEL006663 PE=4 SV=1
CCG028333.1	1236	272	1454	9.45612283	41.12405698	2.120661916	Up	5.36E-13	1.38E-12	AAEL000091-PA OS=Aedes aegypti GN=AAEL000091 PE=4 SV=1
CCG010022.1	2298	178	1455	3.32837805	22.13415696	2.733383211	Up	1.15E-13	3.58E-13	AAEL012263-PA (Fragment) OS=Aedes aegypti GN=AAEL012263 PE=4 SV=1
CCG011952.1	1347	106	1456	3.3814342	37.78712025	3.482187389	Up	0	0	AAEL004699-PA OS=Aedes aegypti GN=AAEL004699 PE=4 SV=1
CCG003119.1	1527	462	1456	13.0006663	33.33284282	1.358358803	Up	0	0	AAEL003001-PA OS=Aedes aegypti GN=AAEL003001 PE=4 SV=1
CCG025473.1	3184	312	1458	4.21060214	16.00790438	1.926685989	Up	9.28E-14	2.93E-13	AAEL004237-PA OS=Aedes aegypti GN=AAEL004237 PE=4 SV=1
CCG007822.1	1734	82	1459	2.03201744	29.41414407	3.855525367	Up	0	0	AAEL008535-PA OS=Aedes aegypti GN=AAEL008535 PE=4 SV=1
CCG003095.1	609	4830	1459	340.794447	83.75061709	-2.024730102	Down	0	0	AAEL000138-PA OS=Aedes aegypti GN=AAEL000138 PE=4 SV=1
CCG015308.1	1630	385	1460	10.1492931	31.31232153	1.625351222	Up	7.93E-13	1.99E-12	AAEL012346-PA OS=Aedes aegypti GN=AAEL012348 PE=4 SV=1
CCG019545.1	1824	401	1460	9.44674538	27.981954	1.566607431	Up	1.05E-13	3.28E-13	AAEL005037-PA OS=Aedes aegypti GN=AAEL005037 PE=3 SV=1
CCG005427.1	1815	408	1460	9.65931224	28.12070749	1.541640516	Up	1.01E-12	2.49E-12	AAEL000088-PA OS=Aedes aegypti GN=AAEL000088 PE=4 SV=1
CCG008924.1	996	472	1460	20.3631674	51.24406033	1.331422808	Up	0	0	AAEL000574-PA OS=Aedes aegypti GN=AAEL000574 PE=4 SV=1
CCG024422.1	1981	449	1461	9.73922809	25.7819497	1.404482032	Up	1.06E-12	2.60E-12	AAEL003526-PA OS=Aedes aegypti GN=AAEL003528 PE=4 SV=1

CCG018632.1	771	9	1462	0.50159223	66.28923559	7.046115798	Up	1.09E-06	2.01E-06	AAEL008678-PA OS=Aedes aegypti GN=AAEL008678 PE=4 SV=1
CCG011419.1	1202	355	1462	12.6907286	42.51996726	1.744365585	Up	2.28E-13	6.52E-13	AAEL010514-PA OS=Aedes aegypti GN=AAEL010514 PE=4 SV=1
CCG028315.1	1350	3220	1462	102.490774	37.85851899	-1.436804173	Down	8.54E-239	5.23E-238	AAEL005699-PA OS=Aedes aegypti GN=AAEL005699 PE=4 SV=1
CCG002370.1	3585	301	1463	3.60777967	14.26609733	1.983407581	Up	0	0	AAEL003327-PA OS=Aedes aegypti GN=AAEL003327 PE=4 SV=1
CCG005171.1	2538	243	1464	4.11412353	20.165058	2.293200538	Up	1.53E-14	5.41E-14	AAEL009285-PA OS=Aedes aegypti GN=AAEL009285 PE=3 SV=1
CCG025892.1	2354	209	1465	3.8150699	21.75610683	2.511639021	Up	0	0	AAEL010641-PA OS=Aedes aegypti GN=AAEL010641 PE=4 SV=1
CCG010746.1	1041	67	1466	2.7655833	49.23038785	4.153893401	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011214 PE=4 SV=1
CCG014394.1	3143	237	1466	3.24016134	16.30570593	2.331239343	Up	2.32E-13	6.61E-13	AAEL005837-PA OS=Aedes aegypti GN=AAEL005837 PE=4 SV=1
CCG006528.1	928	437	1466	20.2346703	55.22503637	1.448493122	Up	5.07E-13	1.31E-12	-
CCG001206.1	2830	138	1467	2.09534395	18.12148128	3.112441903	Up	1.49E-13	4.45E-13	-
CCG006952.1	2592	80	1469	1.32622638	19.81238757	3.901003789	Up	0	0	AAEL003070-PA OS=Aedes aegypti GN=AAEL003070 PE=4 SV=1
CCG020164.1	6389	493	1470	3.31571125	8.043303624	1.278469807	Up	6.97E-14	2.28E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001092 PE=4 SV=1
CCG026217.1	1855	169	1471	3.91476288	27.7216308	2.824015299	Up	0	0	AAEL002249-PA OS=Aedes aegypti GN=AAEL002249 PE=4 SV=1
CCG010899.1	1830	385	1471	9.04008078	28.1003416	1.636180099	Up	7.93E-13	1.99E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_10453 PE=4 SV=1
CCG012755.1	4449	164	1472	1.58395965	11.56632578	2.868323155	Up	1.14E-13	3.54E-13	AAEL009871-PA OS=Aedes aegypti GN=AAEL009871 PE=4 SV=1
CCG020872.1	519	191	1472	15.8135247	99.14948634	2.648446332	Up	6.68E-14	2.19E-13	AAEL000173-PA OS=Aedes aegypti GN=AAEL000173 PE=4 SV=1
CCG023241.1	3027	105	1473	1.49052598	17.01141119	3.512609401	Up	0	0	AAEL007630-PA OS=Aedes aegypti GN=AAEL007630 PE=4 SV=1

CCG019599.1	2463	165	1473	2.87860585	20.90683787	2.860532705	Up	1.31E-14	4.66E-14	AAEL007353-PA OS=Aedes aegypti GN=AAEL007353 PE=3 SV=1
CCG011533.1	1881	190	1474	4.34037723	27.39420519	2.657978405	Up	2.69E-13	7.52E-13	AAEL010885-PA OS=Aedes aegypti GN=AAEL010885 PE=4 SV=1
CCG022619.2	2936	308	1474	4.5077242	17.55057901	1.961047472	Up	0	0	AAEL005821-PA OS=Aedes aegypti GN=AAEL005821 PE=4 SV=1
CCG018175.1	2782	3192	1474	49.3024417	18.52210638	-1.412410923	Down	1.08E-230	6.59E-230	AAEL007197-PA OS=Aedes aegypti GN=AAEL007197 PE=4 SV=1
CCG006490.1	255	3971	1475	669.148299	202.2096402	-1.726474203	Down	0	0	AAEL008802-PA OS=Aedes aegypti GN=AAEL008802 PE=4 SV=1
CCG021535.1	1296	2	1476	0.06631132	39.81359299	9.22979021	Up	9.36E-05	0.00015406	AAEL002586-PA OS=Aedes aegypti GN=AAEL002586 PE=4 SV=1
CCG027496.1	2077	165	1476	3.41358027	24.84276192	2.863467996	Up	1.31E-14	4.67E-14	AAEL002115-PA OS=Aedes aegypti GN=AAEL002115 PE=4 SV=1
CCG000806.1	2773	166	1476	2.57229569	18.60743473	2.854750779	Up	1.83E-13	5.32E-13	AAEL010007-PA OS=Aedes aegypti GN=AAEL010007 PE=4 SV=1
CCG019462.1	3881	200	1476	2.21436406	13.29513438	2.58593402	Up	0	0	AAEL002918-PA OS=Aedes aegypti GN=AAEL002918 PE=4 SV=1
CCG008488.1	975	480	1476	21.1543309	52.92145284	1.322899614	Up	5.41E-13	1.39E-12	AAEL004445-PA OS=Aedes aegypti GN=AAEL004445 PE=4 SV=1
CCG026738.1	2472	3974	1476	69.0783679	20.87314584	-1.726585947	Down	0	0	AAEL006323-PA OS=Aedes aegypti GN=AAEL006323 PE=4 SV=1
CCG019198.1	1489	80	1477	2.30864927	34.67654452	3.90883922	Up	0	0	AAEL007757-PA OS=Aedes aegypti GN=AAEL007757 PE=4 SV=1
CCG027286.1	1540	141	1477	3.93424194	33.52816545	3.091215962	Up	0	0	AAEL001001-PA OS=Aedes aegypti GN=AAEL001001 PE=4 SV=1
CCG019328.1	4008	166	1477	1.77968462	12.88257854	2.855727883	Up	1.83E-13	5.32E-13	AAEL007137-PA (Fragment) OS=Aedes aegypti GN=AAEL007137 PE=4 SV=1
CCG022165.1	824	113	1478	5.89269419	62.70428771	3.411564796	Up	1.42E-13	4.26E-13	-
CCG020326.1	1485	457	1479	13.2236826	34.81703121	1.396669186	Up	1.13E-12	2.77E-12	AAEL009339-PA (Fragment) OS=Aedes aegypti GN=AAEL009339 PE=3 SV=1
CCG022701.1	1656	67	1480	1.73850979	31.24290436	4.167605474	Up	0	0	AAEL010782-PA OS=Aedes aegypti GN=AAEL010782 PE=3 SV=1

CCG010967.1	3030	267	1480	3.78644196	17.07532991	2.172998733	Up	5.21E-13	1.35E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000710 PE=4 SV=1
CCG000393.1	2796	391	1480	6.0090008	18.50438112	1.622669867	Up	9.02E-13	2.25E-12	AAEL010387-PA OS=Aedes aegypti GN=AAEL010387 PE=4 SV=1
CCG004884.2	1785	404	1481	9.7253629	29.00459826	1.576457646	Up	1.18E-13	3.65E-13	AAEL001398-PA (Fragment) OS=Aedes aegypti GN=AAEL001398 PE=4 SV=1
CCG007162.1	1932	2560	1481	56.9371225	26.79772666	-1.087258966	Down	3.50E-123	1.93E-122	-
CCG022070.1	3108	426	1482	5.88967405	16.66929414	1.500933316	Up	4.37E-13	1.15E-12	AAEL004911-PA OS=Aedes aegypti GN=AAEL004911 PE=4 SV=1
CCG012164.1	2690	537	1482	8.57797304	19.25954133	1.166864658	Up	3.67E-13	9.84E-13	AAEL014507-PA OS=Aedes aegypti GN=AAEL014507 PE=4 SV=1
CCG014595.2	3582	225	1484	2.69910393	14.4829935	2.423807389	Up	0	0	AAEL001651-PA OS=Aedes aegypti GN=AAEL001651 PE=4 SV=1
CCG023780.2	592	260	1484	18.8718429	87.6318965	2.215220767	Up	3.22E-13	8.81E-13	AAEL001642-PA OS=Aedes aegypti GN=AAEL001642 PE=4 SV=1
CCG014466.1	1603	496	1484	13.2956883	32.36312085	1.28339227	Up	6.79E-13	1.71E-12	AAEL000858-PA OS=Aedes aegypti GN=AAEL000858 PE=4 SV=1
CCG028508.1	1575	320	1485	8.73035878	32.96066096	1.916632325	Up	0	0	AAEL010960-PA OS=Aedes aegypti GN=AAEL010960 PE=3 SV=1
CCG000866.1	11174	456	1486	1.7535528	4.649006558	1.40664159	Up	1.37E-12	3.31E-12	AAEL013487-PA OS=Aedes aegypti GN=AAEL013487 PE=4 SV=1
CCG018105.1	900	2	1487	0.0954883	57.75884173	9.240502136	Up	9.36E-05	0.00015336	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015628 PE=4 SV=1
CCG006972.1	1725	354	1487	8.81813684	30.13504786	1.772896586	Up	0	0	AAEL010668-PA OS=Aedes aegypti GN=AAEL010668 PE=4 SV=1
CCG000285.1	801	24032	1487	1289.19933	64.89757498	-4.312166962	Down	0	0	AAEL011795-PA OS=Aedes aegypti GN=AAEL011795 PE=4 SV=1
CCG009842.1	3469	440	1489	5.45018254	15.00515253	1.461081529	Up	0	0	AAEL010502-PA OS=Aedes aegypti GN=AAEL010502 PE=4 SV=1
CCG010702.1	1797	475	1489	11.3581658	28.96654096	1.350657539	Up	6.44E-13	1.64E-12	Ultraspiracle OS=Aedes albopictus GN=USP PE=2 SV=1

CCG005150.1	3450	370	1490	4.60834835	15.09792243	1.712028359	Up	0	0	AAEL009668-PA OS=Aedes aegypti GN=AAEL009668 PE=4 SV=1
CCG010891.1	3381	573	1491	7.28235964	15.41638292	1.081986417	Up	5.68E-13	1.46E-12	AAEL001473-PA OS=Aedes aegypti GN=AAEL001473 PE=4 SV=1
CCG018250.1	3924	102	1492	1.11695029	13.29198495	3.572919682	Up	0	0	AGAP004751-PA OS=Anopheles gambiae GN=AGAP004751 PE=4 SV=4
CCG009982.1	2091	199	1492	4.08941998	24.94392585	2.608720404	Up	0	0	AAEL001173-PA OS=Aedes aegypti GN=AAEL001173 PE=4 SV=1
CCG002464.1	4670	291	1492	2.67755734	11.16868286	2.060469681	Up	6.00E-14	2.00E-13	DNA topoisomerase z-binding protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ010004 PE=4 SV=1
CCG000072.1	2104	330	1493	6.73954963	24.80641978	1.87998944	Up	1.76E-13	5.16E-13	P/CAF gamma constitutive coactivator 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005420 PE=4 SV=1
CCG019707.1	1887	488	1493	11.1124698	27.65909233	1.315574316	Up	9.43E-13	2.34E-12	AAEL006589-PA OS=Aedes aegypti GN=AAEL006589 PE=4 SV=1
CCG015232.1	3030	519	1493	7.36016246	17.22531591	1.226720926	Up	0	0	AAEL009264-PA OS=Aedes aegypti GN=AAEL009264 PE=4 SV=1
CCG012022.1	1422	4768	1493	144.078548	36.70373222	-1.972856867	Down	0	0	AAEL001209-PA OS=Aedes aegypti GN=AAEL001209 PE=4 SV=1
CCG013009.1	2298	509	1494	9.51766533	22.72744365	1.255755791	Up	0	0	AAEL006309-PA OS=Aedes aegypti GN=AAEL006309 PE=4 SV=1
CCG009924.1	4495	2690	1494	25.7149246	11.61905795	-1.146112821	Down	9.76E-141	5.49E-140	AAEL003267-PA (Fragment) OS=Aedes aegypti GN=AAEL003267 PE=4 SV=2
CCG008960.1	948	46	1496	2.08502932	55.16622579	4.725645708	Up	3.40E-14	1.18E-13	AAEL009330-PA OS=Aedes aegypti GN=AAEL009330 PE=4 SV=1
CCG018617.1	6055	303	1496	2.15026087	8.637090347	2.00603368	Up	1.37E-13	4.14E-13	DNA-directed RNA polymerase OS=Anopheles gambiae GN=AGAP004703 PE=3 SV=4
CCG005655.1	2199	43	1497	0.84024492	23.79833576	4.823906955	Up	8.46E-14	2.70E-13	AAEL001357-PA OS=Aedes aegypti GN=AAEL001357 PE=4 SV=1

CCG007714.1	2190	217	1497	4.25773169	23.89613714	2.488620477	Up	1.47E-13	4.37E-13	AAEL013357-PA OS=Aedes aegypti GN=AAEL013357 PE=4 SV=1
CCG028365.1	1059	296	1498	12.0104263	49.44995147	2.041681747	Up	2.26E-13	6.47E-13	AAEL008942-PA OS=Aedes aegypti GN=AAEL008942 PE=4 SV=1
CCG004038.1	1313	314	1499	10.2760828	39.91047744	1.957477123	Up	4.97E-14	1.68E-13	AAEL001371-PA OS=Aedes aegypti GN=AAEL001371 PE=4 SV=1
CCG016820.1	2361	278	1500	5.0595452	22.20983277	2.134118916	Up	2.11E-14	7.39E-14	AGAP001943-PA OS=Anopheles gambiae GN=AgaP_AGAP001943 PE=4 SV=5
CCG022929.1	3690	209	1503	2.43378714	14.23910298	2.548583366	Up	0	0	Tyrosine-protein kinase receptor OS=Anopheles gambiae GN=INR PE=3 SV=3
CCG006561.1	1718	234	1503	5.85268795	30.58340512	2.385577778	Up	4.60E-13	1.20E-12	AAEL000142-PA (Fragment) OS=Aedes aegypti GN=AAEL000142 PE=4 SV=1
CCG005168.1	1275	581	1503	19.5807183	41.20963921	1.073548144	Up	0	0	AGAP011626-PA OS=Anopheles gambiae GN=AGAP011626 PE=4 SV=4
CCG009431.1	1017	348	1505	14.703508	51.7327498	1.81491748	Up	0	0	-
CCG024661.1	2507	521	1505	8.929889	20.98612148	1.232721413	Up	0	0	AAEL009877-PA (Fragment) OS=Aedes aegypti GN=AAEL009877 PE=4 SV=1
CCG026363.1	2958	120	1506	1.7431941	17.79823016	3.351928663	Up	0	0	AAEL009529-PA OS=Aedes aegypti GN=AAEL009529 PE=4 SV=1
CCG006599.1	4065	188	1506	1.98728416	12.95133206	2.704230407	Up	1.80E-13	5.25E-13	AAEL008324-PA OS=Aedes aegypti GN=AAEL008324 PE=4 SV=1
CCG016980.1	1642	549	1507	14.3668601	32.08411882	1.159114566	Up	2.32E-13	6.62E-13	AAEL010791-PA OS=Aedes aegypti GN=AAEL010791 PE=4 SV=1
CCG026193.1	2397	125	1508	2.24080802	21.99294175	3.294949632	Up	2.35E-14	8.22E-14	AAEL014106-PA OS=Aedes aegypti GN=AAEL014106 PE=4 SV=1
CCG020759.1	2286	287	1508	5.39471296	23.0608405	2.09582699	Up	2.87E-13	7.98E-13	AAEL013905-PB (Fragment) OS=Aedes aegypti GN=AAEL013905 PE=4 SV=1
CCG020829.1	1725	377	1508	9.39106664	30.56062688	1.702313204	Up	4.66E-13	1.21E-12	AAEL011061-PA OS=Aedes aegypti GN=AAEL011061 PE=4 SV=1

CCG022755.1	894	3556	1508	170.917647	58.96765254	-1.535305692	Down	6.64E-291	4.16E-290	AAEL008485-PA OS=Aedes aegypti GN=AAEL008485 PE=4 SV=1
CCG019407.1	1500	142	1510	4.06780154	35.19133195	3.112898919	Up	2.62E-13	7.34E-13	AAEL002557-PA OS=Aedes aegypti GN=AAEL002557 PE=4 SV=1
CCG009702.1	2987	311	1511	4.47391612	17.68394918	1.982830379	Up	0	0	AAEL010732-PA OS=Aedes aegypti GN=AAEL010732 PE=4 SV=1
CCG001264.1	4130	417	1511	4.33859064	12.7898199	1.559697576	Up	0	0	Connexin43-interacting protein of 150 kDa OS=Culex quinquefasciatus GN=CpipJ_CPIJ015283 PE=4 SV=1
CCG025605.1	1215	77	1512	2.72318483	43.50363332	3.997769087	Up	0	0	AAEL000294-PA OS=Aedes aegypti GN=AAEL000294 PE=4 SV=1
CCG014264.1	1857	170	1512	3.93368599	28.463605	2.855164692	Up	3.53E-13	9.53E-13	AAEL012568-PA OS=Aedes aegypti GN=AAEL012568 PE=4 SV=1
CCG002831.1	2817	558	1512	8.51157682	18.76354792	1.140434316	Up	2.34E-13	6.64E-13	AAEL004227-PA (Fragment) OS=Aedes aegypti GN=AAEL004227 PE=4 SV=1
CCG025879.1	1601	10	1514	0.2683931	33.05860777	6.944534599	Up	5.43E-07	1.01E-06	AAEL015402-PA OS=Aedes aegypti GN=AAEL015402 PE=4 SV=1
CCG012163.1	2127	232	1515	4.68687279	24.8997599	2.409434287	Up	2.91E-13	8.06E-13	AAEL014509-PA OS=Aedes aegypti GN=AAEL014509 PE=4 SV=1
CCG026693.1	2181	272	1516	5.35890317	24.29928821	2.180904401	Up	5.36E-13	1.38E-12	AAEL004419-PA OS=Aedes aegypti GN=AAEL004419 PE=4 SV=1
CCG011587.1	1473	54	1518	1.57526522	36.02624857	4.515381777	Up	0	0	AAEL002569-PA OS=Aedes aegypti GN=AAEL002569 PE=4 SV=1
CCG005481.1	2476	87	1518	1.50984124	21.43241686	3.827325783	Up	0	0	AAEL010595-PA OS=Aedes aegypti GN=AAEL010595 PE=3 SV=1
CCG023476.1	1464	236	1518	6.92681514	36.24772141	2.38762623	Up	4.60E-13	1.20E-12	AAEL005628-PA OS=Aedes aegypti GN=AAEL005628 PE=4 SV=1
CCG017984.1	2753	288	1520	4.49519926	19.30133698	2.102243811	Up	0	0	AAEL004110-PA OS=Aedes aegypti GN=AAEL004110 PE=4 SV=1
CCG026810.1	1767	521	1520	12.6696275	30.07163593	1.24702925	Up	0	0	AAEL012632-PA OS=Aedes aegypti GN=AAEL012632 PE=4 SV=1
CCG010847.1	2443	190	1522	3.34189504	21.77916383	2.704210239	Up	2.69E-13	7.52E-13	AAEL012618-PA OS=Aedes aegypti GN=AAEL012618 PE=4 SV=1

CCG022759.1	5940	471	1523	3.40719613	8.963208001	1.395430181	Up	0	0	AAEL001977-PA OS=Aedes aegypti GN=AAEL001977 PE=4 SV=1
CCG012689.1	2992	263	1524	3.77708563	17.80628804	2.237041402	Up	1.40E-14	4.98E-14	AAEL002238-PA OS=Aedes aegypti GN=AAEL002238 PE=4 SV=1
CCG019724.1	300	4	1525	0.57292979	177.7045736	8.276906731	Up	2.95E-05	5.04E-05	-
CCG010567.3	5802	590	1526	4.36955247	9.194472657	1.073281306	Up	0	0	AAEL011663-PA OS=Aedes aegypti GN=AAEL011663 PE=4 SV=1
CCG016855.3	2314	182	1529	3.37964205	23.09905151	2.772891255	Up	0	0	AAEL014106-PA OS=Aedes aegypti GN=AAEL014106 PE=4 SV=1
CCG015862.1	2310	310	1529	5.76650118	23.13904986	2.00456149	Up	2.14E-13	6.15E-13	AAEL007642-PA OS=Aedes aegypti GN=TUBE PE=4 SV=1
CCG013024.1	7922	463	1529	2.51135914	6.747185709	1.425817512	Up	6.16E-13	1.57E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010714 PE=4 SV=1
CCG013001.1	1404	3397	1529	103.965946	38.07065896	-1.449359608	Down	3.99E-255	2.46E-254	Uncharacterized protein OS=Anopheles darlingi GN=AND_10266 PE=4 SV=1
CCG006089.1	1143	1	1531	0.03759382	46.82512838	10.28257177	Up	0.000587	0.00091592	AAEL005938-PA OS=Aedes aegypti GN=AAEL005938 PE=4 SV=1
CCG000214.1	927	115	1532	5.33065748	57.7735491	3.438023735	Up	6.13E-14	2.03E-13	AAEL003876-PA OS=Aedes aegypti GN=AAEL003876 PE=4 SV=1
CCG024190.2	3441	419	1532	5.23229259	15.56410346	1.572707352	Up	0	0	AAEL001688-PA OS=Aedes aegypti GN=AAEL001688 PE=4 SV=1
CCG021651.1	13993	3269	1532	10.0384523	3.827347961	-1.391119876	Down	8.01E-231	4.88E-230	AAEL013936-PA OS=Aedes aegypti GN=SRPN4 PE=3 SV=2
CCG014660.1	3160	582	1533	7.91404606	16.95918933	1.099579843	Up	1.68E-13	4.94E-13	-
CCG008406.1	1601	337	1534	9.04484732	33.49531328	1.88879119	Up	0	0	AAEL003768-PA OS=Aedes aegypti GN=AAEL003768 PE=4 SV=1
CCG024432.2	2394	504	1535	9.04625992	22.41476811	1.309056221	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_12429 PE=3 SV=1
CCG012084.1	2608	304	1537	5.00874207	20.60232799	2.04028714	Up	0	0	AAEL006016-PA OS=Aedes aegypti GN=AAEL006016 PE=4 SV=1
CCG012350.1	1143	241	1539	9.06011027	47.06980573	2.377201384	Up	2.37E-13	6.72E-13	AAEL000459-PA OS=Aedes aegypti GN=AAEL000459 PE=4 SV=1

CCG017834.1	1278	250	1539	8.40566014	42.09764316	2.324306435	Up	2.10E-13	6.02E-13	Saccharopine dehydrogenase domain-containing protein OS=Aedes aegypti PE=2 SV=1
CCG015977.1	2802	181	1540	2.77570377	19.21332842	2.791181952	Up	1.53E-13	4.55E-13	AAEL003916-PA OS=Aedes aegypti GN=AAEL003916 PE=4 SV=1
CCG007580.1	822	510	1542	26.6600543	65.57866519	1.298546817	Up	0	0	AAEL007528-PA OS=Aedes aegypti GN=AAEL007528 PE=4 SV=1
CCG012283.1	1836	130	1543	3.04251934	29.37942324	3.271467737	Up	0	0	AAEL011201-PA OS=Aedes aegypti GN=AAEL011201 PE=4 SV=1
CCG025411.1	3077	571	1543	7.97390915	17.53026359	1.136488615	Up	0	0	AAEL013977-PA OS=Aedes aegypti GN=AAEL013977 PE=4 SV=1
CCG007669.1	3756	47	1544	0.5376937	14.37049503	4.740181389	Up	0	0	Microcephalin OS=Culex quinquefasciatus GN=CpipJ_CPIJ009302 PE=4 SV=1
CCG012153.1	1880	84	1544	1.91992431	28.71041454	3.902452818	Up	0	0	AAEL009775-PA OS=Aedes aegypti GN=AAEL009775 PE=4 SV=1
CCG016685.1	3342	244	1544	3.13722778	16.15068203	2.364032904	Up	2.78E-13	7.73E-13	AAEL012045-PA (Fragment) OS=Aedes aegypti GN=AAEL012045 PE=4 SV=1
CCG019116.3	2592	301	1544	4.98992674	20.82391178	2.061150564	Up	0	0	AAEL002145-PA OS=Aedes aegypti GN=AAEL002145 PE=4 SV=1
CCG000089.1	742	20076	1544	1162.61508	72.74336838	-3.99841399	Down	0	0	AAEL014505-PA OS=Aedes aegypti GN=AAEL014505 PE=2 SV=1
CCG000612.1	741	0	1545	0.001	72.88871473	16.15340784	Up	4.44E-16	1.65E-15	AAEL001559-PA OS=Aedes aegypti GN=AAEL001559 PE=3 SV=1
CCG023617.1	1371	411	1545	12.8815178	39.39499461	1.612709743	Up	0	0	AAEL013071-PA OS=Aedes aegypti GN=AAEL013071 PE=4 SV=1
CCG020467.1	1028	87	1547	3.63654369	52.60744569	3.854627189	Up	0	0	AAEL014250-PA OS=Aedes aegypti GN=AAEL014250 PE=4 SV=1
CCG003264.1	1952	3485	1547	76.7159452	27.7051507	-1.469372255	Down	4.03E-267	2.50E-266	-
CCG000552.1	2182	214	1548	4.21426361	24.80083063	2.557035974	Up	0	0	AAEL007308-PA (Fragment) OS=Aedes aegypti GN=AAEL007308 PE=4 SV=1
CCG007525.1	462	547	1548	50.8754217	117.1329274	1.203105937	Up	0	0	AAEL007342-PA OS=Aedes aegypti GN=AAEL007342 PE=4 SV=1

CCG024375.1	15493	147	1549	0.40770354	3.49515076	3.099762288	Up	7.02E-14	2.29E-13	AAEL010986-PA (Fragment) OS=Aedes aegypti GN=AAEL010986 PE=3 SV=1
CCG017366.1	3171	220	1549	2.98118625	17.07674889	2.518074919	Up	4.61E-13	1.20E-12	AAEL013086-PA OS=Aedes aegypti GN=AAEL013086 PE=4 SV=1
CCG010641.1	5580	122	1550	0.93948165	9.710632437	3.369628366	Up	8.04E-14	2.58E-13	AAEL008377-PA OS=Aedes aegypti GN=AAEL008377 PE=4 SV=1
CCG002095.1	2633	62	1551	1.01182056	20.59258917	4.347099864	Up	0	0	AAEL006140-PA OS=Aedes aegypti GN=AAEL006140 PE=3 SV=1
CCG027350.1	933	452	1551	20.8170633	58.11391991	1.481117212	Up	1.12E-13	3.49E-13	AAEL003961-PA OS=Aedes aegypti GN=AAEL003961 PE=3 SV=1
CCG021482.1	1947	128	1552	2.82492349	27.86607373	3.302226046	Up	4.15E-14	1.42E-13	AAEL013487-PA OS=Aedes aegypti GN=AAEL013487 PE=4 SV=1
CCG001014.1	2178	138	1553	2.72260026	24.92663169	3.194630861	Up	1.49E-13	4.45E-13	AAEL004489-PA OS=Aedes aegypti GN=AAEL004489 PE=4 SV=1
CCG020084.1	3570	264	1553	3.17759382	15.20734001	2.258761199	Up	0	0	AAEL011147-PA OS=Aedes aegypti GN=AAEL011147 PE=4 SV=1
CCG026664.1	945	72	1554	3.27388454	57.48694403	4.134158991	Up	0	0	AAEL010422-PA OS=Aedes aegypti GN=AAEL010422 PE=4 SV=1
CCG002959.1	3110	357	1554	4.93253867	17.46789778	1.824303728	Up	3.04E-13	8.40E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011015 PE=4 SV=1
CCG020742.1	1728	466	1555	11.587903	31.458403	1.440825924	Up	0	0	AAEL000650-PA OS=Aedes aegypti GN=AAEL000650 PE=4 SV=1
CCG015245.1	2505	412	1556	7.06727771	21.71460226	1.619439022	Up	3.41E-13	9.25E-13	AAEL002694-PA OS=Aedes aegypti GN=AAEL002694 PE=4 SV=2
CCG023719.1	2913	380	1558	5.60538934	18.69721772	1.737937114	Up	0	0	AAEL005469-PA OS=Aedes aegypti GN=AAEL005469 PE=4 SV=1
CCG022407.1	3086	2922	1558	40.6861842	17.64905872	-1.204947741	Down	2.53E-165	1.46E-164	Stranded-chain-amino-acid aminotransferase OS=Culex quinquefasciatus GN=CpipJ_CPIJ015408 PE=3 SV=1
CCG003300.2	1333	332	1559	10.7021395	40.88518641	1.933678985	Up	0	0	Semaphorin-like protein kinase 3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000288 PE=4 SV=1

CCG019292.1	3366	456	1559	5.82121182	16.1913112	1.475828402	Up	1.37E-12	3.31E-12	AAEL007487-PA OS=Aedes aegypti GN=AAEL007487 PE=4 SV=1
CCG011457.1	1242	6363	1559	220.142046	43.88079991	-2.32677299	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000118 PE=4 SV=1
CCG014712.1	3360	393	1560	5.02592432	16.2306285	1.691258015	Up	1.08E-13	3.37E-13	AAEL009261-PA OS=Aedes aegypti GN=AAEL009261 PE=4 SV=1
CCG020026.1	4503	524	1561	5.00025337	12.11855875	1.277145024	Up	0	0	AAEL004621-PA OS=Aedes aegypti GN=AAEL004621 PE=4 SV=1
CCG015908.2	1350	4230	1561	134.638502	40.42212596	-1.735873922	Down	0	0	AAEL007029-PA OS=Aedes aegypti GN=AAEL007029 PE=4 SV=1
CCG021734.1	1389	636	1564	19.6751269	39.36266729	1.000455046	Up	9.12E-13	2.27E-12	AAEL000424-PA OS=Aedes aegypti GN=AAEL000424 PE=3 SV=1
CCG023671.2	7563	8046	1564	45.7139342	7.229240364	-2.660718022	Down	0	0	AAEL002070-PA (Fragment) OS=Aedes aegypti GN=AAEL002070 PE=4 SV=1
CCG027497.1	1718	66	1565	1.65075814	31.84499601	4.269866026	Up	0	0	AAEL002104-PA OS=Aedes aegypti GN=AAEL002104 PE=4 SV=1
CCG005885.1	2817	4186	1565	63.8520799	19.42126487	-1.717096451	Down	0	0	AAEL012110-PA OS=Aedes aegypti GN=AAEL012110 PE=4 SV=1
CCG024581.1	1320	89	1566	2.8972018	41.47322835	3.83944827	Up	0	0	AAEL006279-PA OS=Aedes aegypti GN=AAEL006279 PE=4 SV=1
CCG012546.1	2226	125	1566	2.41294556	24.59328905	3.349397416	Up	2.35E-14	8.23E-14	GA19507, isoform A OS=Drosophila pseudoobscura pseudoobscura GN=GA19507 PE=4 SV=1
CCG021745.1	1883	222	1568	5.06600164	29.11023791	2.522607182	Up	0	0	AAEL008112-PA OS=Aedes aegypti GN=AAEL008112 PE=4 SV=1
CCG026695.1	4517	503	1568	4.78498484	12.13517334	1.342608458	Up	1.17E-13	3.62E-13	AAEL004431-PA OS=Aedes aegypti GN=AAEL004431 PE=4 SV=1
CCG004634.1	2892	178	1569	2.64474853	18.96595306	2.84220941	Up	1.15E-13	3.57E-13	AAEL002098-PA (Fragment) OS=Aedes aegypti GN=AAEL002098 PE=4 SV=1
CCG005065.1	1202	208	1569	7.43569451	45.63189372	2.617503123	Up	3.93E-13	1.05E-12	AAEL013463-PA OS=Aedes aegypti GN=AAEL013463 PE=4 SV=1

CCG022456.1	3432	252	1569	3.15512037	15.98179961	2.340662917	Up	5.42E-14	1.81E-13	Structural maintenance of chromosomes protein OS=Aedes aegypti GN=AAEL001655 PE=3 SV=1
CCG010489.1	1428	363	1569	10.9229788	38.41003939	1.814117103	Up	0	0	AAEL013291-PA OS=Aedes aegypti GN=AAEL013291 PE=4 SV=1
CCG007646.1	942	411	1569	18.7479415	58.22668392	1.634948257	Up	0	0	AAEL001264-PA OS=Aedes aegypti GN=AAEL001264 PE=4 SV=1
CCG011911.1	1061	113	1570	4.57641848	51.72902406	3.498683085	Up	1.42E-13	4.26E-13	Proliferating cell nuclear antigen OS=Aedes aegypti GN=PCNA PE=2 SV=1
CCG000471.1	5316	141	1571	1.13971644	10.33097306	3.180229317	Up	0	0	AAEL002319-PA OS=Aedes aegypti GN=AAEL002319 PE=4 SV=1
CCG003044.1	1846	245	1571	5.70291711	29.75051615	2.38314273	Up	3.40E-13	9.23E-13	AAEL007211-PA OS=Aedes aegypti GN=AAEL007211 PE=4 SV=1
CCG014495.1	5502	117	1572	0.91375117	9.988079078	3.450333986	Up	9.44E-14	2.97E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005250 PE=4 SV=1
CCG020280.1	4541	408	1575	3.86074691	12.12492533	1.651023975	Up	1.01E-12	2.49E-12	AAEL008475-PA OS=Aedes aegypti GN=AAEL008475 PE=4 SV=1
CCG007917.1	1395	3250	1575	100.108701	39.46902216	-1.342774686	Down	2.06E-217	1.24E-216	AAEL006866-PA OS=Aedes aegypti GN=AAEL006866 PE=4 SV=1
CCG024297.1	674	9557	1576	609.290436	81.74220207	-2.89797708	Down	0	0	-
CCG020286.1	1775	279	1577	6.75411603	31.05870562	2.201158837	Up	0	0	AAEL009618-PA OS=Aedes aegypti GN=AAEL009618 PE=4 SV=1
CCG002531.1	3431	522	1577	6.53751136	16.06796924	1.297374152	Up	0	0	AAEL013627-PA OS=Aedes aegypti GN=AAEL013627 PE=4 SV=1
CCG011283.1	4581	3297	1577	30.9258273	12.03431619	-1.361658023	Down	2.67E-225	1.62E-224	AAEL003348-PA OS=Aedes aegypti GN=AAEL003348 PE=4 SV=2
CCG013007.1	4539	359	1578	3.39857562	12.15337315	1.83835466	Up	0	0	AAEL006320-PA OS=Aedes aegypti GN=AAEL006320 PE=4 SV=1
CCG028391.1	1314	3481	1578	113.833825	41.98185749	-1.439091405	Down	1.34E-258	8.27E-258	AAEL000044-PB OS=Aedes aegypti GN=AAEL000044 PE=3 SV=1
CCG001914.1	4968	138	1581	1.19360374	11.12500716	3.220410399	Up	1.49E-13	4.45E-13	AAEL004377-PA OS=Aedes aegypti GN=AAEL004377 PE=4 SV=1

CCG012271.1	1923	194	1581	4.33496023	28.74104814	2.729022014	Up	5.97E-13	1.53E-12	AAEL012611-PA OS=Aedes aegypti GN=AAEL012611 PE=4 SV=1
CCG005196.2	7444	290	1582	1.67399557	7.429338239	2.149937998	Up	1.62E-13	4.78E-13	AAEL001977-PA OS=Aedes aegypti GN=AAEL001977 PE=4 SV=1
CCG009639.1	1008	373	1582	15.900507	54.86507327	1.786815268	Up	3.38E-13	9.19E-13	AAEL012282-PA OS=Aedes aegypti GN=AAEL012282 PE=4 SV=1
CCG014576.2	1344	228	1583	7.28950855	41.17481557	2.49786873	Up	1.05E-13	3.28E-13	AAEL012608-PA OS=Aedes aegypti GN=AAEL012608 PE=4 SV=1
CCG010434.1	1605	264	1583	7.06791896	34.4790979	2.286364625	Up	0	0	AAEL001111-PA OS=Aedes aegypti GN=AAEL001111 PE=4 SV=1
CCG019620.1	4187	97	1584	0.99547749	13.22519952	3.731756982	Up	3.62E-14	1.25E-13	AAEL003361-PA OS=Aedes aegypti GN=AAEL003361 PE=4 SV=1
CCG024188.1	1293	109	1584	3.62235195	42.82591679	3.563485499	Up	0	0	AAEL001681-PA OS=Aedes aegypti GN=AAEL001681 PE=4 SV=1
CCG005594.1	4259	197	1584	1.98756462	13.00162254	2.709618004	Up	0	0	AAEL010595-PA OS=Aedes aegypti GN=AAEL010595 PE=3 SV=1
CCG004453.1	796	273	1584	14.7371075	69.56521408	2.238912683	Up	1.53E-14	5.41E-14	Small nuclear ribonucleoprotein-associated protein OS=Aedes aegypti GN=AAEL002377 PE=2 SV=1
CCG006775.1	2529	162	1585	2.75250969	21.90939845	2.992730326	Up	0	0	AAEL007521-PA OS=Aedes aegypti GN=AAEL007521 PE=4 SV=1
CCG012493.1	811	278	1585	14.7294528	68.32166299	2.213639256	Up	2.11E-14	7.38E-14	AAEL004436-PA OS=Aedes aegypti GN=AAEL004436 PE=4 SV=1
CCG011684.1	675	4701	1586	299.260329	82.1390029	-1.865261706	Down	0	0	AAEL001865-PA OS=Aedes aegypti GN=AAEL001865 PE=4 SV=1
CCG013361.1	1395	12	1588	0.36963213	39.79479822	6.7503459	Up	1.30E-07	2.49E-07	AAEL000584-PA OS=Aedes aegypti GN=AAEL000584 PE=4 SV=1
CCG005938.1	2721	320	1588	5.05340503	20.40196381	2.013380306	Up	0	0	AAEL009475-PA (Fragment) OS=Aedes aegypti GN=AAEL009475 PE=3 SV=1
CCG026514.1	2196	5499	1588	107.600442	25.27948247	-2.08964517	Down	0	0	AAEL010004-PA OS=Aedes aegypti GN=AAEL010004 PE=4 SV=1
CCG009753.1	1382	159	1589	4.94369595	40.19442966	3.023333658	Up	5.53E-14	1.84E-13	AAEL006189-PA OS=Aedes aegypti GN=AAEL006189 PE=4 SV=1

CCG026812.1	1351	413	1589	13.1358256	41.11672968	1.646218642	Up	5.88E-13	1.51E-12	AAEL013394-PA OS=Aedes aegypti GN=AAEL013394 PE=3 SV=1
CCG025528.2	4578	3863	1589	36.2586467	12.13383613	-1.579289349	Down	0	0	AAEL011573-PA OS=Aedes aegypti GN=AAEL011573 PE=4 SV=1
CCG014903.1	1719	8	1591	0.1999755	32.35521719	7.338031324	Up	2.19E-06	3.97E-06	AAEL011516-PA OS=Aedes aegypti GN=AAEL011516 PE=4 SV=1
CCG017798.1	1509	191	1591	5.43884646	36.85793131	2.760602496	Up	6.68E-14	2.19E-13	AAEL001492-PA OS=Aedes aegypti GN=AAEL001492 PE=4 SV=1
CCG028653.1	2386	370	1592	6.66337041	23.32505307	1.807556364	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_14317 PE=4 SV=1
CCG017361.1	1533	423	1592	11.8566195	36.30370295	1.614423971	Up	1.18E-13	3.65E-13	AGAP001896-PA OS=Anopheles gambiae GN=AGAP001896 PE=4 SV=1
CCG007356.1	4710	2785	1592	25.4077942	11.81604599	-1.104523788	Down	2.85E-137	1.60E-136	AAEL000597-PA OS=Aedes aegypti GN=AAEL000597 PE=3 SV=1
CCG015725.1	1644	1	1593	0.02613731	33.87380468	10.33984376	Up	0.000587	0.0009178	AAEL000889-PA (Fragment) OS=Aedes aegypti GN=AAEL000889 PE=4 SV=2
CCG023596.1	1464	180	1593	5.28316409	38.03861673	2.847990659	Up	3.32E-13	9.05E-13	AAEL003563-PA OS=Aedes aegypti GN=AAEL003563 PE=4 SV=1
CCG027778.1	1191	91	1594	3.28316192	46.78714792	3.832954478	Up	0	0	AAEL013795-PA OS=Aedes aegypti GN=AAEL013795 PE=4 SV=1
CCG007892.1	1368	271	1594	8.5122793	40.73354764	2.258600076	Up	0	0	AAEL002753-PA OS=Aedes aegypti GN=AAEL002753 PE=4 SV=1
CCG004168.2	2019	3959	1595	84.2581374	27.6168655	-1.60926644	Down	0	0	AAEL013171-PA OS=Aedes aegypti GN=HPX2 PE=4 SV=1
CCG012857.1	2608	2969	1596	48.9176158	21.39317858	-1.193203238	Down	2.27E-165	1.31E-164	AAEL002938-PA OS=Aedes aegypti GN=AAEL002938 PE=3 SV=1
CCG028221.1	887	343	1598	16.6162559	62.98007473	1.922300131	Up	0	0	-
CCG028230.1	3411	629	1600	7.92376519	16.39790174	1.049253187	Up	6.78E-13	1.71E-12	AAEL012553-PA OS=Aedes aegypti GN=HOP PE=4 SV=2
CCG000651.1	314	9567	1600	1309.20844	178.1313466	-2.877681489	Down	0	0	AAEL010673-PA OS=Aedes aegypti GN=AAEL010673 PE=4 SV=1

CCG001040.1	4413	251	1601	2.44400711	12.68257446	2.375527242	Up	7.22E-14	2.35E-13	AAEL001012-PA OS=Aedes aegypti GN=AAEL001612 PE=4 SV=1
CCG024802.1	1569	327	1602	8.95545138	35.6935369	1.994824811	Up	6.36E-13	1.62E-12	AAEL013210-PA OS=Aedes aegypti GN=AAEL013210 PE=2 SV=1
CCG022336.1	1824	22	1603	0.51827531	30.72265223	5.889440295	Up	8.38E-11	1.80E-10	AAEL015126-PA OS=Aedes aegypti GN=AAEL015128 PE=4 SV=1
CCG003630.1	2009	175	1603	3.74300824	27.89353791	2.897660802	Up	3.55E-13	9.58E-13	AAEL000990-PA OS=Aedes aegypti GN=AAEL000990 PE=4 SV=1
CCG026605.1	1779	216	1604	5.21723591	31.51943561	2.594884128	Up	1.38E-13	4.17E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006272 PE=4 SV=1
CCG008302.1	1098	2	1605	0.0782691	51.10021331	9.350670786	Up	9.36E-05	0.00015395	AAEL010638-PA OS=Aedes aegypti GN=AAEL010638 PE=3 SV=1
CCG017448.1	1022	632	1606	26.5722821	54.93443493	1.047788633	Up	0	0	AAEL013212-PA OS=Aedes aegypti GN=AAEL013212 PE=4 SV=1
CCG020460.1	1071	437	1608	17.5329356	52.48637633	1.581875425	Up	5.07E-13	1.31E-12	AAEL011462-PA OS=Aedes aegypti GN=AAEL011462 PE=4 SV=1
CCG013292.1	1794	68	1609	1.62873019	31.35332627	4.266798973	Up	0	0	AAEL012962-PA OS=Aedes aegypti GN=AAEL012962 PE=3 SV=1
CCG006104.1	3624	292	1609	3.46224131	15.52093469	2.164437256	Up	0	0	AAEL001436-PA OS=Aedes aegypti GN=AAEL001436 PE=4 SV=1
CCG025210.1	506	489	1609	41.5260874	111.1617931	1.42057116	Up	1.91E-13	5.51E-13	AAEL005832-PA OS=Aedes aegypti GN=AAEL005832 PE=4 SV=1
CCG014847.1	1569	611	1609	16.7332746	35.84950116	1.099233245	Up	2.21E-13	6.33E-13	AAEL010303-PA OS=Aedes aegypti GN=AAEL010303 PE=4 SV=1
CCG000353.1	1896	7578	1609	171.742958	29.66659669	-2.53333961	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_22820 PE=3 SV=1
CCG025308.1	2784	363	1610	5.60273479	20.21653219	1.851332439	Up	0	0	AAEL006677-PA OS=Aedes aegypti GN=AAEL006677 PE=4 SV=1
CCG000590.1	5292	521	1611	4.23039148	10.64206045	1.33091442	Up	0	0	AAEL008579-PA OS=Aedes aegypti GN=AAEL008579 PE=4 SV=1
CCG025666.1	3182	620	1611	8.37248129	17.69886357	1.079929577	Up	4.74E-13	1.23E-12	AAEL012070-PA OS=Aedes aegypti GN=AAEL012070 PE=4 SV=1

CCG007504.1	1518	361	1612	10.2187577	37.12301855	1.861094205	Up	1.35E-13	4.08E-13	AGAP000765-PA OS=Anopheles gambiae GN=AgaP_AGAP000765 PE=4 SV=3
CCG027545.1	648	7742	1612	513.38223	86.96410827	-2.56154136	Down	0	0	AAEL0008072-PA OS=Aedes aegypti GN=AAEL0008072 PE=3 SV=1
CCG012410.1	1996	243	1613	5.23128533	28.25035092	2.433031423	Up	1.53E-14	5.41E-14	AAEL0006306-PA OS=Aedes aegypti GN=AAEL006306 PE=4 SV=1
CCG021995.1	2147	531	1613	10.627354	26.26348413	1.305275876	Up	2.40E-13	6.77E-13	AAEL014644-PA OS=Aedes aegypti GN=AAEL014644 PE=4 SV=1
CCG008969.2	2928	21093	1613	309.549389	19.25809441	-4.006632753	Down	0	0	Ubiquitin carboxyl-terminal hydrolase OS=Aedes aegypti GN=AAEL010854 PE=3 SV=1
CCG012348.1	2349	471	1614	8.61589826	24.01986322	1.479154817	Up	0	0	AAEL000440-PA OS=Aedes aegypti GN=AAEL000440 PE=4 SV=1
CCG011310.1	1704	639	1614	16.1136505	33.11188891	1.039065946	Up	4.28E-13	1.13E-12	AAEL000321-PA OS=Aedes aegypti GN=AAEL000321 PE=4 SV=1
CCG019458.1	5013	190	1615	1.62861551	11.26224157	2.789776045	Up	2.69E-13	7.52E-13	AAEL009806-PA (Fragment) OS=Aedes aegypti GN=AAEL009806 PE=4 SV=1
CCG001397.1	3602	420	1615	5.01035218	15.67396363	1.645386136	Up	1.84E-13	5.32E-13	AAEL0005272-PA OS=Aedes aegypti GN=AAEL0005272 PE=4 SV=1
CCG020877.1	2013	405	1617	8.6451776	28.08123872	1.69963907	Up	2.34E-13	6.64E-13	AAEL000199-PA OS=Aedes aegypti GN=AAEL000199 PE=3 SV=1
CCG023331.1	12801	470	1617	1.57767169	4.415868568	1.484900221	Up	1.05E-12	2.59E-12	AAEL011578-PA OS=Aedes aegypti GN=AAEL011578 PE=4 SV=1
CCG002865.1	3124	540	1617	7.42754696	18.09460101	1.28460157	Up	9.26E-13	2.31E-12	AAEL009495-PA (Fragment) OS=Aedes aegypti GN=AAEL009495 PE=4 SV=1
CCG023805.1	1284	201	1619	6.72657061	44.07901098	2.712148783	Up	0	0	AAEL003101-PA OS=Aedes aegypti GN=AAEL003101 PE=4 SV=1
CCG006076.1	3286	339	1620	4.43297019	17.23445173	1.958949838	Up	1.30E-13	3.98E-13	AAEL007048-PA OS=Aedes aegypti GN=AAEL007048 PE=4 SV=1
CCG004050.1	2206	394	1620	7.67455822	25.67198929	1.742039482	Up	5.49E-13	1.41E-12	AAEL009424-PA OS=Aedes aegypti GN=AAEL009424 PE=4 SV=1

CCG018181.1	1935	182	1621	4.04159778	29.2854608	2.857186939	Up	0	0	AAEL002594-PA (Fragment) OS=Aedes aegypti GN=AAEL002594 PE=4 SV=1
CCG017031.1	2904	441	1621	6.52536259	19.51355601	1.580346734	Up	0	0	AAEL003791-PA OS=Aedes aegypti GN=AAEL003791 PE=4 SV=1
CCG016689.2	4089	244	1622	2.56410253	13.8670396	2.43513397	Up	2.78E-13	7.73E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014927 PE=4 SV=1
CCG016831.1	2910	425	1622	6.27564853	19.48533503	1.634552277	Up	1.58E-12	3.80E-12	AAEL001144-PA OS=Aedes aegypti GN=AAEL001144 PE=4 SV=1
CCG018290.1	3894	16066	1622	177.286019	14.56145992	-3.605851853	Down	0	0	Malic enzyme OS=Aedes aegypti GN=AAEL001091 PE=3 SV=1
CCG026511.1	1493	318	1623	9.15229444	38.00219906	2.053877533	Up	1.20E-13	3.69E-13	AAEL010893-PA OS=Aedes aegypti GN=AAEL010893 PE=2 SV=1
CCG026054.2	628	273	1624	18.6795184	90.4016584	2.27489198	Up	1.53E-14	5.41E-14	AAEL012710-PA OS=Aedes aegypti GN=AAEL012710 PE=4 SV=1
CCG005889.1	1065	481	1624	19.4069881	53.30726899	1.457756037	Up	0	0	Zinc-finger protein MCG4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006296 PE=4 SV=1
CCG021881.1	2810	245	1626	3.74647152	20.22852599	2.432786807	Up	3.40E-13	9.24E-13	AAEL012247-PA OS=Aedes aegypti GN=AAEL012247 PE=4 SV=1
CCG024919.1	4842	490	1626	4.34844485	11.73939654	1.432786807	Up	0	0	AAEL001583-PA OS=Aedes aegypti GN=AAEL001583 PE=4 SV=1
CCG011813.1	4457	650	1626	6.26662048	12.75345704	1.025128838	Up	0	0	AAEL013098-PA OS=Aedes aegypti GN=AAEL013098 PE=4 SV=1
CCG003094.1	2146	143	1627	2.8633141	26.50378206	3.210440403	Up	0	0	AAEL000136-PA OS=Aedes aegypti GN=AAEL000136 PE=4 SV=1
CCG009489.1	1227	7	1628	0.24514111	46.38310887	7.563843266	Up	4.31E-06	7.71E-06	AAEL007584-PA OS=Aedes aegypti GN=AAEL007584 PE=4 SV=1
CCG004973.1	3138	224	1628	3.06731056	18.13641638	2.563843266	Up	2.04E-13	5.88E-13	Structural maintenance of chromosomes protein OS=Aedes aegypti GN=AAEL001655 PE=3 SV=1
CCG008516.1	2367	67	1629	1.21629583	24.05873801	4.305994902	Up	0	0	AAEL004244-PA OS=Aedes aegypti GN=AAEL004244 PE=4 SV=1

CCG017539.1	12302	2677	1629	9.35051045	4.629087373	-1.014317331	Down	5.19E-115	2.81E-114	AAEL001969-PA OS=Aedes aegypti GN=AAEL001969 PE=4 SV=1
CCG026798.1	3204	469	1630	6.28988937	17.78464143	1.49952534	Up	0	0	AGAP002411-PA OS=Anopheles gambiae GN=AgaP_AGAP002411 PE=4 SV=5
CCG021072.1	1893	3123	1630	70.8898474	30.10142163	-1.235747401	Down	1.04E-183	6.11E-183	Catalase (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG023752.1	1080	2	1631	0.07957358	52.79347168	9.373854271	Up	9.36E-05	0.00015343	AAEL009263-PA OS=Aedes aegypti GN=AAEL009263 PE=2 SV=1
CCG013205.1	3895	218	1632	2.4049813	14.64747309	2.606554221	Up	0	0	PRP16 mRNA splicing factor ATP-dependent RNA helicase PRP16 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003160 PE=4 SV=1
CCG006974.1	330	563	1633	73.3089715	172.9905029	1.238631167	Up	2.33E-14	8.15E-14	AAEL010652-PA OS=Aedes aegypti GN=AAEL010652 PE=4 SV=1
CCG020640.1	1896	404	1634	9.1559983	30.12754443	1.718293989	Up	1.18E-13	3.64E-13	AAEL014054-PA OS=Aedes aegypti GN=AAEL014054 PE=4 SV=1
CCG017735.1	16950	3010	1634	7.63061364	3.37001913	-1.1790423	Down	1.73E-164	1.00E-163	-
CCG023819.1	1896	102	1635	2.31166294	30.14598234	3.704962782	Up	0	0	KanBP-type and CSHC4-type zinc finger-containing protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004842 PE=4 SV=1
CCG025661.1	1946	390	1636	8.61161177	29.38938376	1.770939923	Up	6.59E-13	1.67E-12	AAEL001350-PA OS=Aedes aegypti GN=AAEL001350 PE=4 SV=1
CCG015414.1	6354	595	1636	4.02376331	9.000903494	1.161524379	Up	7.20E-13	1.81E-12	AAEL013079-PA (Fragment) OS=Aedes aegypti GN=AAEL013079 PE=4 SV=1
CCG021164.1	2220	398	1637	7.70358305	25.77779238	1.74252719	Up	7.64E-13	1.92E-12	AAEL010933-PA OS=Aedes aegypti GN=AAEL010933 PE=4 SV=1
CCG016901.1	1739	249	1638	6.1526532	32.92792257	2.420030913	Up	0	0	AAEL004838-PA OS=Aedes aegypti GN=AAEL004838 PE=4 SV=1
CCG026839.1	642	10464	1638	700.366516	89.1926127	-2.97311398	Down	0	0	-
CCG018432.1	2580	152	1639	2.53155026	22.20799055	3.132985829	Up	8.39E-14	2.69E-13	AAEL011815-PA OS=Aedes aegypti GN=AAEL011815 PE=4 SV=1

CCG022140.1	13455	8392	1639	26.8005955	4.258388378	-2.65388562	Down	0	0	AAEL009869-PA OS=Aedes aegypti GN=AAEL009869 PE=4 SV=1
CCG020840.1	50895	646	1641	0.5454062	1.127154577	1.047282373	Up	1.08E-12	2.64E-12	AAEL002063-PA OS=Aedes aegypti GN=AAEL002063 PE=4 SV=1
CCG018533.1	1023	394	1642	16.5494384	56.11093887	1.761499796	Up	5.49E-13	1.41E-12	AAEL008696-PA (Fragment) OS=Aedes aegypti GN=AAEL008696 PE=4 SV=1
CCG018318.2	1470	56	1644	1.63694227	39.09619525	4.577952865	Up	0	0	AAEL013830-PA OS=Aedes aegypti GN=AAEL013830 PE=4 SV=1
CCG015802.1	3201	104	1644	1.3960801	17.954204	3.684868069	Up	0	0	Nkt1, nikkomycin biosynthesis protein P6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005325 PE=4 SV=1
CCG021985.1	2514	106	1644	1.81177083	22.86054376	3.657387333	Up	0	0	AAEL012420-PA OS=Aedes aegypti GN=AAEL012420 PE=4 SV=1
CCG013842.1	17163	6334	1645	15.8579677	3.350601019	-2.24271608	Down	0	0	Sensory appendage protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002629 PE=4 SV=1
CCG026209.1	4890	607	1646	5.33387094	11.76714183	1.141509118	Up	0	0	Purative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002903 PE=4 SV=1
CCG013250.1	3909	4203	1646	46.2015335	14.7202158	-1.650141917	Down	0	0	AAEL000882-PA OS=Aedes aegypti GN=AAEL008882 PE=4 SV=1
CCG010716.1	1917	91	1647	2.03977353	30.03457582	3.880143403	Up	0	0	AAEL014679-PA OS=Aedes aegypti GN=AAEL014679 PE=4 SV=1
CCG001477.1	981	314	1647	13.7538192	58.6914188	2.093317295	Up	4.97E-14	1.68E-13	AAEL001496-PA OS=Aedes aegypti GN=AAEL001496 PE=4 SV=1
CCG003854.1	1412	3082	1648	93.7908797	40.80116156	-1.200837416	Down	2.51E-173	1.46E-172	AAEL0006976-PA OS=Aedes aegypti GN=AAEL006976 PE=4 SV=1
CCG024813.1	1698	242	1649	6.1240729	33.94946902	2.47082565	Up	3.87E-13	1.03E-12	AAEL0008220-PA OS=Aedes aegypti GN=AAEL008220 PE=4 SV=1
CCG010820.1	3832	583	1649	6.53741004	15.04337119	1.202336814	Up	2.70E-13	7.54E-13	AAEL002437-PA OS=Aedes aegypti GN=AAEL002437 PE=4 SV=1
CCG002037.1	1956	161	1650	3.53687488	29.4893439	3.059646635	Up	0	0	AAEL000834-PA OS=Aedes aegypti GN=AAEL008834 PE=4 SV=1

CCG014756.1	1681	454	1650	11.6051514	34.31359707	1.564015026	Up	3.16E-13	8.66E-13	AAEL002636-PA OS=Aedes aegypti GN=AAEL002636 PE=2 SV=1
CCG018422.1	1025	354	1651	14.8402791	56.30840483	1.923832059	Up	0	0	AAEL007015-PA OS=Aedes aegypti GN=AAEL007015 PE=4 SV=1
CCG014866.1	1752	181	1653	4.43922487	32.98289469	2.893338326	Up	1.53E-13	4.55E-13	AAEL014400-PA OS=Aedes aegypti GN=AAEL014400 PE=4 SV=1
CCG019678.1	795	3042	1653	164.420041	72.68683208	-1.177620225	Down	6.76E-166	3.91E-165	ATP synthase subunit beta OS=Culex quinquefasciatus GN=CpipJ_CPIJ006209 PE=3 SV=1
CCG016808.1	3948	206	1655	2.24208848	14.65449546	2.708428178	Up	1.10E-13	3.43E-13	AAEL007436-PA (Fragment) OS=Aedes aegypti GN=AAEL007436 PE=4 SV=1
CCG012214.1	1496	580	1655	16.6593891	38.67376207	1.215019615	Up	7.73E-14	2.49E-13	AAEL008007-PA OS=Aedes aegypti GN=AAEL008007 PE=4 SV=1
CCG011788.1	3081	345	1656	4.81160612	18.78964827	1.96534761	Up	9.59E-13	2.38E-12	AAEL012378-PA OS=Aedes aegypti GN=AAEL012378 PE=4 SV=1
CCG007773.2	3120	569	1657	7.83646763	18.56598225	1.244386249	Up	0	0	Rab1-interacting OS=Culex quinquefasciatus GN=CpipJ_CPIJ006771 PE=4 SV=1
CCG003605.1	1106	4120	1659	160.068089	52.43741516	-1.610017247	Down	0	0	Peptidyl-prolyl cis-trans isomerase OS=Aedes aegypti GN=AAEL013279 PE=2 SV=1
CCG001648.1	1257	3778	1661	129.148494	46.19387249	-1.483257425	Down	1.68E-293	1.06E-292	AAEL007881-PA OS=Aedes aegypti GN=AAEL007881 PE=4 SV=1
CCG028239.1	765	154	1662	8.65011651	75.94856993	3.13423133	Up	1.82E-13	5.28E-13	AAEL008784-PA OS=Aedes aegypti GN=AAEL008784 PE=3 SV=1
CCG014769.1	1359	356	1662	11.2562366	42.75250625	1.92528444	Up	7.93E-13	1.99E-12	AAEL012501-PA OS=Aedes aegypti GN=AAEL012501 PE=4 SV=1
CCG027156.1	1203	360	1662	12.8587734	48.29647215	1.909164774	Up	5.08E-13	1.32E-12	AAEL003508-PA OS=Aedes aegypti GN=AAEL003508 PE=3 SV=1
CCG022136.1	1671	214	1663	5.50300611	34.79091219	2.660418671	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018768 PE=4 SV=1
CCG023740.2	3360	439	1663	5.61420044	17.30226615	1.623808528	Up	0	0	AAEL012019-PA OS=Aedes aegypti GN=AAEL012019 PE=4 SV=1

CCG022735.1	2586	76	1664	1.26283829	22.49442094	4.154825409	Up	0	0	AAEL007007-PA OS=Aedes aegypti GN=AAEL007007 PE=3 SV=1
CCG022364.1	944	52	1665	2.36697691	61.6584013	4.703179948	Up	0	0	Protease III zinc metalloprotease OS=Culex quinquefasciatus GN=CpipJ_CPIJ001046 PE=4 SV=1
CCG021197.1	2074	9487	1665	196.554422	28.06438323	-2.808116566	Down	0	0	AAEL0006833-PA OS=Aedes aegypti GN=AAEL006833 PE=2 SV=1
CCG007758.1	1812	259	1666	6.14192123	32.14155028	2.387677601	Up	0	0	AAEL010781-PA OS=Aedes aegypti GN=AAEL010781 PE=4 SV=1
CCG026015.1	1461	4732	1666	139.173706	39.86344223	-1.803748469	Down	0	0	AAEL011126-PA OS=Aedes aegypti GN=AAEL011126 PE=3 SV=1
CCG001083.1	3273	94	1667	1.23408343	17.80490296	3.850762741	Up	0	0	AAEL013313-PA OS=Aedes aegypti GN=AAEL013313 PE=4 SV=1
CCG026089.1	1567	581	1668	15.931982	37.21149053	1.223822424	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_00721 PE=4 SV=1
CCG024511.1	1284	253	1669	8.46677793	45.44031459	2.424087868	Up	3.13E-13	8.61E-13	AAEL002891-PA OS=Aedes aegypti GN=AAEL002891 PE=4 SV=1
CCG025935.2	2672	563	1669	9.05387746	21.8358398	1.270090331	Up	2.33E-14	8.15E-14	AAEL006172-PA OS=Aedes aegypti GN=AAEL006172 PE=4 SV=1
CCG010378.2	4313	104	1670	1.0361355	13.53589664	3.707505873	Up	0	0	DNA polymerase OS=Aedes aegypti GN=AAEL009851 PE=3 SV=1
CCG016407.1	2502	320	1671	5.49572945	23.34743425	2.086881127	Up	0	0	AAEL008020-PA OS=Aedes aegypti GN=AAEL008020 PE=4 SV=1
CCG023563.1	855	35057	1671	1761.85963	68.32196548	-4.688605701	Down	0	0	AAEL002759-PA OS=Aedes aegypti GN=AAEL002759 PE=3 SV=1
CCG019455.1	2280	134	1672	2.52541423	25.63606963	3.343583145	Up	0	0	AAEL011626-PA OS=Aedes aegypti GN=AAEL011626 PE=4 SV=1
CCG010968.1	897	186	1672	8.91011219	65.16191612	2.870513525	Up	6.43E-13	1.64E-12	AAEL000198-PA OS=Aedes aegypti GN=AAEL000198 PE=4 SV=1
CCG020627.1	3299	244	1673	3.1781192	17.72815915	2.479797596	Up	2.78E-13	7.73E-13	Zinc finger protein 363 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000911 PE=4 SV=1
CCG020554.1	1131	9813	1674	372.822286	51.74195872	-2.849081528	Down	0	0	AAEL005964-PA OS=Aedes aegypti GN=AAEL005964 PE=3 SV=1

CCG009235.1	1935	199	1675	4.41910966	30.26104062	2.775633964	Up	0	0	AAEL000674-PA OS=Aedes aegypti GN=AAEL000674 PE=4 SV=1
CCG007076.1	1041	411	1675	16.9649961	56.24890835	1.729264	Up	0	0	AAEL004448-PA OS=Aedes aegypti GN=AAEL004448 PE=4 SV=1
CCG024810.1	1948	75	1676	1.6543789	30.07703895	4.184300947	Up	0	0	AAEL014528-PA OS=Aedes aegypti GN=AAEL014528 PE=4 SV=1
CCG010546.1	3705	198	1676	2.29635829	15.81378458	2.783763017	Up	0	0	AAEL012637-PA OS=Aedes aegypti GN=AAEL012637 PE=4 SV=1
CCG012790.1	1461	3841	1677	112.968344	40.12664623	-1.493286071	Down	2.24E-301	1.42E-300	AAEL007452-PA OS=Aedes aegypti GN=AAEL007452 PE=4 SV=1
CCG025513.1	1172	145	1678	5.31622143	50.05118466	3.234931114	Up	0	0	AAEL009879-PA OS=Aedes aegypti GN=AAEL009879 PE=4 SV=1
CCG002758.1	1914	203	1678	4.5573961	30.64785184	2.749504287	Up	9.28E-14	2.93E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016156 PE=4 SV=1
CCG011767.1	12830	529	1682	1.77170613	4.582994663	1.371151282	Up	0	0	AAEL011810-PA OS=Aedes aegypti GN=AAEL011810 PE=4 SV=1
CCG026745.1	1673	140	1684	3.59579369	35.18812796	3.29070661	Up	0	0	AAEL000001-PA OS=Aedes aegypti GN=AAEL000001 PE=4 SV=1
CCG026501.1	2157	670	1684	13.3471127	27.2924145	1.031972342	Up	8.60E-13	2.15E-12	AAEL010063-PA OS=Aedes aegypti GN=AAEL010063 PE=3 SV=1
CCG024261.1	2431	157	1685	2.77509187	24.23064433	3.126225331	Up	0	0	Elongation factor 1-alpha OS=Culex quinquefasciatus GN=CpipJ_CPIJ002376 PE=4 SV=1
CCG024910.1	4281	143	1687	1.43533568	13.7758965	3.262686125	Up	0	0	AAEL011477-PA OS=Aedes aegypti GN=AAEL011477 PE=4 SV=1
CCG011908.1	11679	441	1687	1.62254071	5.049628642	1.637922617	Up	0	0	AGAP000420-PA OS=Anopheles gambiae GN=AgaP_AGAP000420 PE=4 SV=3
CCG015357.1	2637	490	1687	7.98451648	22.36428249	1.485919523	Up	0	0	AAEL011277-PA (Fragment) OS=Aedes aegypti GN=AAEL011277 PE=4 SV=1
CCG013353.1	4527	54	1689	0.51256145	13.04275005	4.669379314	Up	0	0	DNA polymerase OS=Aedes aegypti GN=AAEL008247 PE=3 SV=1

CCG016140.1	2098	78	1690	1.59754018	28.15990836	4.139718516	Up	0	0	AAEL015455-PA OS=Aedes aegypti GN=AAEL015455 PE=4 SV=1
CCG011097.1	1399	593	1690	18.2137617	42.22979825	1.21323244	Up	1.13E-12	2.78E-12	AAEL005197-PA OS=Aedes aegypti GN=AAEL005197 PE=4 SV=1
CCG021451.2	1463	542	1696	15.9190678	40.52579454	1.348084617	Up	0	0	AAEL005550-PA OS=Aedes aegypti GN=AAEL003550 PE=2 SV=1
CCG006465.1	7287	5401	1696	31.8484337	8.136302649	-1.968777175	Down	0	0	AAEL012612-PA OS=Aedes aegypti GN=AAEL012612 PE=4 SV=1
CCG019148.1	34442	499	1697	0.62255088	1.722437596	1.468188048	Up	0	0	AAEL005267-PA OS=Aedes aegypti GN=AAEL005267 PE=4 SV=1
CCG027349.1	1715	653	1697	16.361071	34.59136774	1.080144872	Up	5.96E-13	1.53E-12	Dehydrogenase-like protein OS=Aedes aegypti PE=2 SV=1
CCG011415.1	2807	396	1698	6.0619932	21.14683077	1.802577327	Up	4.78E-13	1.24E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010150 PE=4 SV=1
CCG028219.1	1569	454	1698	12.4335625	37.83247544	1.60538546	Up	3.16E-13	8.65E-13	AAEL010254-PA OS=Aedes aegypti GN=AAEL010254 PE=4 SV=1
CCG003451.1	2110	121	1699	2.46414118	28.14886836	3.513920104	Up	0	0	AAEL006112-PA OS=Aedes aegypti GN=AAEL006112 PE=4 SV=1
CCG011101.1	1326	335	1699	10.855853	44.79193985	2.044766056	Up	8.48E-14	2.70E-13	AAEL005208-PA OS=Aedes aegypti GN=AAEL005208 PE=2 SV=1
CCG000445.1	3600	424	1699	5.06087985	16.49836451	1.704862886	Up	0	0	AAEL011751-PA OS=Aedes aegypti GN=AAEL011751 PE=4 SV=1
CCG009700.1	1449	264	1700	7.82885434	41.01385129	2.389238115	Up	0	0	AAEL005958-PA OS=Aedes aegypti GN=AAEL003938 PE=4 SV=1
CCG007082.1	1548	44	1701	1.22136197	38.41345529	4.975049011	Up	3.91E-14	1.34E-13	-
CCG020830.1	2728	280	1701	4.41038332	21.79766451	2.305197613	Up	0	0	AAEL011062-PA OS=Aedes aegypti GN=AAEL011062 PE=4 SV=1
CCG020871.1	3429	79	1702	0.98997056	17.35170226	4.131547777	Up	0	0	AAEL000201-PA (Fragment) OS=Aedes aegypti GN=AAEL000201 PE=4 SV=1
CCG016503.1	1086	293	1702	11.5931236	54.78728091	2.240571671	Up	4.19E-13	1.11E-12	AAEL013484-PA OS=Aedes aegypti GN=AAEL013484 PE=4 SV=1
CCG021716.1	1988	470	1702	10.1588407	29.92906794	1.558811579	Up	1.05E-12	2.59E-12	AAEL005037-PA OS=Aedes aegypti GN=AAEL005037 PE=3 SV=1

CCG002247.1	530	535	1703	43.3751095	112.3281988	1.372780842	Up	2.42E-14	8.43E-14	AAEL004278-PA OS=Aedes aegypti GN=AAEL004278 PE=4 SV=1
CCG028345.1	3690	156	1704	1.81660667	16.14333431	3.151620605	Up	1.42E-13	4.26E-13	AAEL009515-PA (Fragment) OS=Aedes aegypti GN=AAEL009515 PE=4 SV=1
CCG017868.1	2111	256	1705	5.21091997	28.23489431	2.437869228	Up	0	0	AAEL004920-PA OS=Aedes aegypti GN=AAEL004920 PE=4 SV=1
CCG017181.1	2318	119	1706	2.20595273	25.72856781	3.543897372	Up	0	0	AAEL013590-PA OS=Aedes aegypti GN=AAEL013590 PE=4 SV=1
CCG013126.2	1426	586	1706	17.6579695	41.82245454	1.243958281	Up	0	0	AAEL001525-PA (Fragment) OS=Aedes aegypti GN=AAEL001525 PE=4 SV=1
CCG024183.1	1808	285	1709	6.77343715	33.0440791	2.286431776	Up	4.57E-13	1.20E-12	AAEL013982-PA OS=Aedes aegypti GN=AAEL013982 PE=4 SV=1
CCG004593.1	4184	389	1710	3.99503508	14.28744103	1.838467469	Up	0	0	DNA repair protein RAD50 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004115 PE=4 SV=1
CCG014899.1	1458	595	1710	17.5356599	41.00044807	1.225347955	Up	7.20E-13	1.81E-12	AAEL012117-PA (Fragment) OS=Aedes aegypti GN=AAEL012117 PE=3 SV=1
CCG001453.2	1989	369	1711	7.97176072	30.07220289	1.915460242	Up	0	0	GE11757 OS=Drosophila yakuba GN=GE11757 PE=4 SV=1
CCG007767.1	2100	96	1712	1.96433073	28.49931897	3.858817689	Up	0	0	AGAP013200-PA OS=Anopheles gambiae GN=AgaP_AGAP013200 PE=4 SV=1
CCG002143.1	1095	465	1712	18.2474215	54.65622816	1.582693284	Up	0	0	AAEL005358-PA OS=Aedes aegypti GN=AAEL005358 PE=4 SV=1
CCG020922.1	2013	218	1713	4.65345362	29.74839946	2.676438315	Up	0	0	Zinc finger protein 250 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001374 PE=4 SV=1
CCG015205.2	282	394	1713	60.0357285	212.3529366	1.82257082	Up	5.49E-13	1.41E-12	Putative secreted peptide OS=Aedes albopictus PE=4 SV=1
CCG011322.1	2169	100	1714	1.98108504	27.62493609	3.801608408	Up	0	0	AAEL001915-PA OS=Aedes aegypti GN=AAEL001915 PE=3 SV=1

CCG003051.1	3724	388	1715	4.47697557	16.09920641	1.846393223	Up	3.43E-13	9.27E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001573 PE=4 SV=1
CCG026288.1	4038	301	1718	3.20304362	14.87328368	2.215207848	Up	0	0	AAEL001716-PA OS=Aedes aegypti GN=AAEL001716 PE=4 SV=1
CCG013989.1	1625	502	1718	13.2743426	36.95896584	1.477283971	Up	6.14E-13	1.57E-12	AAEL013749-PA OS=Aedes aegypti GN=AAEL013749 PE=4 SV=1
CCG011969.1	3412	334	1720	4.20629876	17.62257797	2.066801761	Up	3.15E-13	8.64E-13	AAEL003075-PA OS=Aedes aegypti GN=AAEL003075 PE=4 SV=1
CCG005757.1	1919	421	1720	9.42691937	31.33310894	1.73282963	Up	4.57E-13	1.20E-12	AAEL009276-PA OS=Aedes aegypti GN=AAEL009276 PE=4 SV=1
CCG015865.1	3495	533	1722	6.55303821	17.224078	1.394190908	Up	0	0	AAEL007639-PA OS=Aedes aegypti GN=AAEL007639 PE=4 SV=1
CCG004091.1	1884	644	1725	14.6881683	32.00797634	1.123776972	Up	2.78E-13	7.74E-13	AAEL010521-PA OS=Aedes aegypti GN=AAEL010521 PE=4 SV=1
CCG016878.1	4347	496	1726	4.90291888	13.88037398	1.501333643	Up	6.79E-13	1.71E-12	AAEL005532-PA (Fragment) OS=Aedes aegypti GN=AAEL005532 PE=4 SV=1
CCG017519.1	1842	208	1727	4.85217416	32.77575678	2.755925853	Up	3.93E-13	1.05E-12	AAEL005943-PA OS=Aedes aegypti GN=AAEL005943 PE=4 SV=1
CCG010634.1	852	320	1727	16.1388675	70.86026289	2.134437476	Up	0	0	AAEL010053-PA OS=Aedes aegypti GN=AAEL010053 PE=4 SV=1
CCG026825.1	2100	487	1727	9.96488607	28.74902095	1.528587609	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_09750 PE=4 SV=1
CCG014566.1	1622	597	1727	15.8156175	37.22129716	1.23477845	Up	0	0	AAEL011995-PA OS=Aedes aegypti GN=AAEL011995 PE=4 SV=1
CCG002974.1	9534	682	1727	3.07377376	6.332383468	1.042737642	Up	1.08E-12	2.66E-12	Ubiquitin carboxyl-terminal hydrolase OS=Aedes aegypti GN=AAEL009702 PE=3 SV=1
CCG009746.1	2077	74	1728	1.53093903	29.08420908	4.24774734	Up	0	0	AAEL003361-PA OS=Aedes aegypti GN=AAEL003361 PE=4 SV=1
CCG018113.1	2486	7097	1728	122.669431	24.29923663	-2.335792885	Down	0	0	AAEL003746-PA OS=Aedes aegypti GN=AAEL003746 PE=4 SV=1

CCG004939.1	1209	11139	1728	395.897332	49.9651797	-2.986131395	Down	0	0	AAEL012950-PA OS=Aedes aegypti GN=AAEL012950 PE=2 SV=1
CCG004955.1	1632	258	1729	6.79300952	37.03606651	2.446808102	Up	1.69E-13	4.97E-13	AAEL007853-PA OS=Aedes aegypti GN=AAEL007853 PE=4 SV=1
CCG002662.1	1701	556	1733	14.0453689	35.61592807	1.34242807	Up	1.85E-13	5.35E-13	AAEL001200-PA OS=Aedes aegypti GN=AAEL001200 PE=3 SV=1
CCG012519.1	3492	565	1733	6.95243415	17.34899589	1.319262086	Up	4.06E-13	1.08E-12	AGAP003818-PB OS=Anopheles gambiae GN=AgaP_AGAP003818 PE=4 SV=1
CCG018101.1	33362	589	1733	0.75862279	1.815919119	1.259245319	Up	0	0	AAEL011390-PA OS=Aedes aegypti GN=AAEL011390 PE=4 SV=1
CCG007630.1	3371	129	1735	1.6443476	17.99246817	3.451805896	Up	0	0	AAEL007480-PA (Fragment) OS=Aedes aegypti GN=AAEL007480 PE=4 SV=1
CCG027230.1	525	285	1736	23.3264274	115.5953685	2.309046327	Up	4.57E-13	1.20E-12	AAEL014943-PA OS=Aedes aegypti GN=AAEL014943 PE=4 SV=1
CCG013826.1	2433	100	1738	1.76612144	24.97225032	3.821669381	Up	0	0	AAEL011879-PA OS=Aedes aegypti GN=AAEL011879 PE=4 SV=1
CCG012262.1	921	380	1739	17.7290979	66.00699599	1.896499813	Up	0	0	AAEL010906-PA OS=Aedes aegypti GN=AAEL010906 PE=4 SV=1
CCG013298.1	795	39	1740	2.10794925	76.51245482	5.181782576	Up	7.33E-14	2.38E-13	AAEL010906-PA OS=Culex quinquefasciatus GN=CpipJ_CPIJ000938 PE=4 SV=1
CCG001759.1	2235	478	1740	9.18994772	27.21583963	1.566317987	Up	5.04E-13	1.31E-12	AAEL004520-PA OS=Aedes aegypti GN=AAEL004520 PE=4 SV=1
CCG007419.1	2704	565	1741	8.97851333	22.50826918	1.325906634	Up	4.06E-13	1.08E-12	AAEL004110-PA OS=Aedes aegypti GN=AAEL004110 PE=4 SV=1
CCG016698.1	1296	89	1742	2.95085369	46.9886714	3.993108681	Up	0	0	AAEL009227-PA OS=Aedes aegypti GN=AAEL009227 PE=3 SV=1
CCG008381.1	1320	430	1742	13.9977166	46.13433192	1.720649263	Up	6.22E-13	1.59E-12	AAEL012752-PA OS=Aedes aegypti GN=AAEL012752 PE=4 SV=1
CCG015037.2	10342	428	1743	1.77828722	5.89173046	1.728203072	Up	1.52E-13	4.51E-13	-
CCG003851.1	1209	288	1745	10.2359666	50.45673529	2.301399523	Up	0	0	AAEL006836-PA OS=Aedes aegypti GN=AAEL006836 PE=2 SV=1

CCG001220.1	11940	3513	1746	12.6426028	5.111989216	-1.306336814	Down	4.47E-225	2.71E-224	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009271 PE=4 SV=1
CCG000794.1	1598	617	1747	16.5909426	38.21784075	1.203850418	Up	4.51E-13	1.18E-12	Malate dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ018614 PE=4 SV=1
CCG023640.1	1568	3836	1748	105.122386	38.97134426	-1.431584332	Down	1.62E-282	1.01E-281	ATP synthase subunit gamma OS=Aedes aegypti GN=AAEL008848 PE=3 SV=1
CCG028282.1	2847	430	1749	6.4899845	21.47594874	1.726434928	Up	6.22E-13	1.59E-12	AAEL010111-PA OS=Aedes aegypti GN=AAEL010111 PE=4 SV=1
CCG022187.1	3559	686	1749	8.28244955	17.17955214	1.052563012	Up	2.38E-13	6.73E-13	AAEL009416-PA OS=Aedes aegypti GN=AAEL009416 PE=4 SV=1
CCG011972.1	2028	14950	1749	316.764069	30.14892804	-3.393230086	Down	0	0	AAEL003425-PB (Fragment) OS=Aedes aegypti GN=AAEL003425 PE=4 SV=1
CCG027734.1	1713	298	1750	7.47517858	35.71335922	2.25628389	Up	3.92E-13	1.05E-12	AAEL009883-PA OS=Aedes aegypti GN=AAEL009883 PE=4 SV=1
CCG027486.1	3855	438	1750	4.8821644	15.86951604	1.700665351	Up	4.14E-13	1.10E-12	AAEL000212-PA OS=Aedes aegypti GN=AAEL000212 PE=4 SV=1
CCG019420.2	2106	311	1752	6.34548313	29.0820992	2.196329493	Up	0	0	Kar OS=Culex quinquefasciatus GN=CpipJ_CPIJ004499 PE=3 SV=1
CCG011135.1	1761	349	1754	8.5158645	34.81931713	2.03166301	Up	4.55E-14	1.54E-13	Lysine--tRNA ligase OS=Aedes aegypti GN=AAEL014702 PE=3 SV=1
CCG016409.1	1401	414	1755	12.6976946	43.79141737	1.786081562	Up	8.70E-13	2.17E-12	AGAP009746-PA OS=Anopheles gambiae GN=AGAP009746 PE=4 SV=2
CCG020870.1	1650	230	1756	5.98972058	37.20408122	2.634900282	Up	3.85E-13	1.03E-12	AAEL000170-PA OS=Aedes aegypti GN=AAEL000170 PE=4 SV=1
CCG016805.1	4626	296	1756	2.74946854	13.26993818	2.270936968	Up	2.26E-13	6.47E-13	AAEL007447-PA OS=Aedes aegypti GN=AAEL007447 PE=4 SV=2
CCG014301.3	4932	356	1759	3.10162723	12.467885	2.00711954	Up	7.93E-13	1.99E-12	AAEL006529-PA OS=Aedes aegypti GN=AAEL006529 PE=4 SV=2
CCG026122.1	485	5	1760	0.44298695	126.8589013	8.161744822	Up	1.59E-05	2.76E-05	-

CCG020285.1	1860	460	1763	10.6269236	33.13518384	1.640639912	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009258 PE=4 SV=1
CCG020231.1	1359	254	1764	8.03113509	45.37630627	2.498263363	Up	2.35E-13	6.67E-13	AAEL000424-PA OS=Aedes aegypti GN=AAEL000424 PE=3 SV=1
CCG024654.1	2334	275	1764	5.06284362	26.42090841	2.383660241	Up	4.31E-13	1.14E-12	AAEL010336-PA OS=Aedes aegypti GN=AAEL010336 PE=4 SV=1
CCG017516.1	933	11203	1765	515.959203	66.13221705	-2.963831824	Down	0	0	AAEL001022-PA OS=Aedes aegypti GN=AAEL001022 PE=4 SV=1
CCG019274.1	327	13895	1765	1825.88521	188.6891697	-3.274512542	Down	0	0	AAEL004457-PA OS=Aedes aegypti GN=AAEL004457 PE=3 SV=1
CCG014340.1	2691	136	1766	2.17164025	22.94177509	3.40111999	Up	1.70E-13	4.99E-13	Unif OS=Culex quinquefasciatus GN=CpipJ_CPIJ006150 PE=4 SV=1
CCG001257.1	4474	415	1766	3.98579344	13.79890853	1.791615305	Up	0	0	AAEL008683-PA OS=Aedes aegypti GN=AAEL008683 PE=4 SV=1
CCG015711.1	2891	110	1767	1.6349605	21.36675028	3.708039815	Up	0	0	AAEL007007-PA OS=Aedes aegypti GN=AAEL007007 PE=3 SV=1
CCG023534.1	3051	368	1767	5.18284573	20.24623896	1.965837572	Up	3.69E-13	9.90E-13	AAEL014249-PA OS=Aedes aegypti GN=AAEL014249 PE=4 SV=1
CCG017958.1	3621	225	1768	2.67003322	17.06882997	2.676434572	Up	0	0	AAEL010795-PA OS=Aedes aegypti GN=AAEL010795 PE=4 SV=1
CCG002173.1	832	187	1769	9.65786102	74.3283553	2.944137077	Up	3.31E-13	9.03E-13	-
CCG016886.1	531	438	1769	35.4439619	116.4617544	1.716244477	Up	4.14E-13	1.10E-12	AAEL015613-PA OS=Aedes aegypti GN=AAEL015613 PE=4 SV=1
CCG005849.1	2787	179	1770	2.75980714	22.2017043	3.008031072	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_12961 PE=4 SV=1
CCG013386.1	1850	659	1770	15.3065163	33.44656751	1.127712194	Up	0	0	-
CCG021042.1	1706	321	1771	8.08516108	36.29021581	2.166232213	Up	0	0	AAEL001971-PA OS=Aedes aegypti GN=AAEL001971 PE=4 SV=1
CCG019820.1	3756	577	1771	6.60104816	16.48325563	1.320234192	Up	4.04E-13	1.07E-12	AAEL009294-PA OS=Aedes aegypti GN=AAEL009294 PE=4 SV=1
CCG017877.1	639	6	1774	0.40347169	97.05161658	7.910140997	Up	8.38E-06	1.48E-05	-
CCG014350.1	2031	488	1774	10.3245842	30.53470359	1.564366161	Up	9.43E-13	2.34E-12	AAEL012446-PA OS=Aedes aegypti GN=IAP6 PE=4 SV=2

CCG008229.1	1464	45	1775	1.32079102	42.38452273	5.004063417	Up	0	0	AAEL003156-PA OS=Aedes aegypti GN=AAEL003156 PE=4 SV=1
CCG003486.1	2604	77	1775	1.27061043	23.82908651	4.229129972	Up	0	0	AAEL004075-PA OS=Aedes aegypti GN=AAEL004075 PE=4 SV=1
CCG022217.1	2325	475	1775	8.77876299	26.68857689	1.60413281	Up	6.44E-13	1.64E-12	AAEL014992-PA (Fragment) OS=Aedes aegypti GN=AAEL014992 PE=4 SV=1
CCG022190.1	2213	272	1776	5.28141338	28.05508339	2.409266229	Up	5.36E-13	1.38E-12	AAEL007869-PA OS=Aedes aegypti GN=AAEL007869 PE=4 SV=1
CCG028366.1	3838	317	1776	3.5490896	16.17662833	2.18839004	Up	2.22E-13	6.35E-13	AAEL004918-PA OS=Aedes aegypti GN=AAEL004918 PE=4 SV=1
CCG019283.1	6000	497	1776	3.55932635	10.34764992	1.539627029	Up	7.04E-14	2.29E-13	AAEL001647-PA (Fragment) OS=Aedes aegypti GN=AAEL001647 PE=4 SV=1
CCG012358.1	1212	537	1776	19.0385705	51.22598973	1.427950792	Up	3.67E-13	9.84E-13	Forkhead box subgroup O OS=Aedes aegypti GN=FoxO PE=2 SV=1
CCG003518.3	2679	209	1779	3.3522488	23.21417483	2.791804867	Up	0	0	DNA mismatch repair protein spellchecker 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017719 PE=3 SV=1
CCG010179.1	1315	260	1779	8.49591711	47.29336455	2.476796186	Up	3.22E-13	8.81E-13	AAEL006458-PA OS=Aedes aegypti GN=AAEL006458 PE=3 SV=1
CCG016012.1	1314	98	1780	3.20474429	47.35596092	3.885264886	Up	0	0	Salivary serpin putative anticoagulant OS=Aedes albopictus PE=2 SV=1
CCG005133.1	2651	258	1780	4.18189043	23.47255098	2.488747474	Up	1.69E-13	4.97E-13	AAEL006317-PA OS=Aedes aegypti GN=AAEL006317 PE=4 SV=1
CCG003248.1	1413	555	1780	16.8777089	44.03802736	1.383630768	Up	0	0	AAEL006449-PA OS=Aedes aegypti GN=AAEL006449 PE=4 SV=1

CCG027197.1	369	5	1782	0.58224573	168.8228976	8.17966673	Up	1.59E-05	2.76E-05	Chromosome undetermined SCAF14565, whole genome shotgun sequence OS=Tetraodon nigroviridis GN=GSTENG00016696001 PE=3 SV=1
CCG002656.1	2261	220	1783	4.1810445	27.56771671	2.721044478	Up	4.61E-13	1.20E-12	AAEL009486-PA OS=Aedes aegypti GN=AAEL009486 PE=4 SV=1
CCG005081.1	2637	376	1783	6.12689428	23.63693875	1.94781534	Up	3.71E-14	1.28E-13	AAEL014423-PA (Fragment) OS=Aedes aegypti GN=AAEL014423 PE=4 SV=1
CCG022352.1	2839	501	1783	7.58289434	21.95512768	1.533737398	Up	0	0	AAEL001472-PA OS=Aedes aegypti GN=AAEL001472 PE=3 SV=1
CCG026883.1	2490	642	1783	11.0789436	25.03237248	1.175974704	Up	0	0	AAEL006155-PA OS=Aedes aegypti GN=AAEL006155 PE=4 SV=1
CCG013811.1	2061	725	1783	15.1155059	30.24289543	1.000567007	Up	0	0	AAEL012307-PA OS=Aedes aegypti GN=AAEL012307 PE=4 SV=1
CCG019325.1	1303	85	1784	2.8030909	47.86305891	4.093822168	Up	0	0	Gag-like protein OS=Aedes aegypti PE=4 SV=1
CCG013830.1	816	7133	1785	375.616565	76.47123044	-2.296271695	Down	0	0	mitochondrial cytochrome c1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014410 PE=4 SV=1
CCG009823.1	10263	721	1786	3.01872539	6.083550844	1.01097412	Up	0	0	AAEL007594-PA OS=Aedes aegypti GN=AAEL007594 PE=3 SV=1
CCG022420.1	3864	415	1787	4.61502067	16.16729829	1.808669597	Up	0	0	Alto guanine nucleotide exchange factor OS=Culex quinquefasciatus GN=CpipJ_CPIJ015585 PE=4 SV=1
CCG009798.1	6454	3556	1787	23.6752985	9.679336937	-1.290402486	Down	1.65E-223	9.98E-223	Growth factor receptor-bound protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002888 PE=4 SV=1
CCG018536.1	1587	388	1788	10.505518	39.38588461	1.906531383	Up	3.43E-13	9.27E-13	Dolichyl-diphosphooligosaccharide protein glycotransferase OS=Aedes aegypti PE=2 SV=1
CCG000120.1	26358	4102	1792	6.68722404	2.376706578	-1.492443651	Down	0	0	AAEL006530-PA OS=Aedes aegypti GN=AAEL006530 PE=4 SV=1

CCG028294.1	1881	595	1793	13.592234	33.32280184	1.293727118	Up	7.20E-13	1.81E-12	AAEL009890-PA OS=Aedes aegypti GN=AAEL009890 PE=4 SV=1
CCG003328.1	1769	179	1795	4.34798332	35.47207847	3.028265555	Up	0	0	AAEL010525-PA OS=Aedes aegypti GN=AAEL010525 PE=4 SV=1
CCG006804.1	2280	204	1797	3.84466047	27.55264182	2.841262555	Up	6.59E-14	2.17E-13	AGAP001909-PA OS=Anopheles gambiae GN=AgaP_AGAP001909 PE=4 SV=5
CCG023205.1	2853	240	1797	3.61469902	22.01893563	2.606797302	Up	0	0	AAEL003572-PA OS=Aedes aegypti GN=AAEL003572 PE=4 SV=1
CCG001308.1	9346	257	1798	1.18159874	6.725335078	2.50886596	Up	4.14E-13	1.10E-12	TXR1 DNA-binding protein 45 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014288 PE=4 SV=1
CCG002092.1	2727	273	1798	4.3017006	23.04913151	2.421733368	Up	1.53E-14	5.42E-14	AAEL014827-PA OS=Aedes aegypti GN=AAEL014827 PE=4 SV=1
CCG006233.1	1290	191	1799	6.36218551	48.75189141	2.937863847	Up	6.68E-14	2.19E-13	AAEL009913-PA OS=Aedes aegypti GN=AAEL009913 PE=4 SV=1
CCG018240.2	4537	299	1800	2.83181632	13.86927445	2.292092721	Up	4.80E-13	1.25E-12	AAEL001387-PA OS=Aedes aegypti GN=AAEL001387 PE=4 SV=1
CCG010208.1	5885	303	1800	2.21237546	10.6924211	2.272920412	Up	1.37E-13	4.14E-13	AAEL001719-PA OS=Aedes aegypti GN=AAEL001719 PE=4 SV=1
CCG022477.1	1598	450	1800	12.1003633	39.37728297	1.702313204	Up	9.36E-13	2.32E-12	AGAP008634-PA (Fragment) OS=Anopheles gambiae GN=AGAP008634 PE=4 SV=4
CCG006212.1	4737	106	1801	0.96153512	13.29108222	3.788975215	Up	0	0	AAEL013242-PA OS=Aedes aegypti GN=AAEL013242 PE=4 SV=1
CCG002729.1	822	625	1801	32.6716352	76.59349935	1.22918329	Up	0	0	AAEL004515-PA OS=Aedes aegypti GN=AAEL004515 PE=4 SV=1
CCG015777.1	1989	221	1805	4.77441496	31.72432859	2.732193766	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006272 PE=4 SV=1
CCG024898.1	4590	5539	1805	51.8538911	13.74720906	-1.915313498	Down	0	0	AAEL013600-PA (Fragment) OS=Aedes aegypti GN=AAEL013600 PE=3 SV=1
CCG019294.1	1740	111	1806	2.74117273	36.28428037	3.726479515	Up	0	0	Strain OS=Culex quinquefasciatus GN=CpipJ_CPIJ013090 PE=4 SV=1

CCG014528.1	1665	209	1808	5.39379852	37.96069934	2.815133034	Up	0	0	Arginine N-methyltransferase OS=Culex quinquefasciatus GN=CpipJ_CPIJ009499 PE=4 SV=1
CCG000744.1	2121	372	1808	7.53641739	29.79941745	1.983333355	Up	1.17E-13	3.60E-13	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC003691 PE=4 SV=1
CCG003639.1	2898	8809	1808	130.614352	21.80971857	-2.582270372	Down	0	0	AAEL002971-PA OS=Aedes aegypti GN=AAEL002971 PE=4 SV=1
CCG027154.1	1613	157	1809	4.18242302	39.20615169	3.228669148	Up	0	0	AAEL001626-PA OS=Aedes aegypti GN=AAEL001626 PE=4 SV=1
CCG000148.1	544	7309	1809	577.32682	116.2491226	-2.312168422	Down	0	0	AAEL009964-PA OS=Aedes aegypti GN=AAEL009964 PE=4 SV=1
CCG003210.1	1599	329	1810	8.84117742	39.57128265	2.162143412	Up	0	0	T-complex protein 1 subunit alpha OS=Aedes aegypti GN=AAEL009320 PE=3 SV=1
CCG006791.1	30382	517	1810	0.73120113	2.082630536	1.510066715	Up	8.39E-14	2.69E-13	AAEL001869-PA OS=Aedes aegypti GN=AAEL001869 PE=4 SV=1
CCG020926.1	2581	143	1811	2.38073307	24.52903496	3.365012698	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000595 PE=4 SV=1
CCG023619.1	2854	380	1811	5.7212681	22.18270471	1.955028426	Up	0	0	Elongation factor 1-alpha OS=Culex quinquefasciatus GN=CpipJ_CPIJ002376 PE=4 SV=1
CCG018046.1	1210	130	1813	4.61658306	52.37963288	3.504108601	Up	0	0	AAEL012185-PA OS=Aedes aegypti GN=AAEL012185 PE=2 SV=1
CCG018640.1	1220	211	1814	7.43165082	51.97894595	2.806172756	Up	1.30E-13	3.97E-13	AAEL004532-PA OS=Aedes aegypti GN=AAEL004532 PE=3 SV=1
CCG022470.1	780	479	1814	26.3878242	81.30040265	1.623390099	Up	0	0	AAEL006931-PA OS=Aedes aegypti GN=AAEL006931 PE=4 SV=1
CCG002497.1	15369	707	1815	1.97668048	4.128393021	1.062500632	Up	0	0	AAEL013105-PA OS=Aedes aegypti GN=AAEL013105 PE=1 SV=1
CCG015852.1	1371	144	1817	4.51323252	46.33055353	3.359730907	Up	7.53E-14	2.44E-13	AAEL000436-PA OS=Aedes aegypti GN=AAEL000436 PE=4 SV=1

CCG021545.1	1639	2	1818	0.05243409	38.77617277	9.530449688	Up	9.36E-05	0.00015375	AAEL002638-PA OS=Aedes aegypti GN=CYP9J6 PE=3 SV=1
CCG013603.1	2313	204	1818	3.78980798	27.47693349	2.858024346	Up	6.59E-14	2.17E-13	AAEL007904-PA OS=Aedes aegypti GN=AAEL007904 PE=4 SV=1
CCG024875.1	2932	139	1819	2.03710543	21.68796229	3.412301959	Up	1.46E-13	4.37E-13	AAEL004869-PA (Fragment) OS=Aedes aegypti GN=AAEL004869 PE=4 SV=1
CCG023099.1	1791	4536	1820	108.827871	35.52432369	-1.615168986	Down	0	0	AAEL001479-PA OS=Aedes aegypti GN=AAEL001479 PE=3 SV=1
CCG023751.1	576	5443	1820	406.049072	110.458444	-1.878150382	Down	0	0	AAEL009257-PA OS=Aedes aegypti GN=AAEL009257 PE=4 SV=1
CCG009138.1	2790	119	1821	1.83275929	22.81685376	3.638010647	Up	0	0	AGAP003524-PA OS=Anopheles gambiae GN=AgaP_AGAP003524 PE=4 SV=5
CCG003615.1	1476	228	1822	6.63760128	43.15310317	2.700730433	Up	1.05E-13	3.28E-13	AAEL006783-PA OS=Aedes aegypti GN=AAEL006783 PE=4 SV=1
CCG027323.1	2223	404	1822	7.80916454	28.65226283	1.875408965	Up	1.18E-13	3.64E-13	AAEL006734-PA OS=Aedes aegypti GN=AAEL006734 PE=3 SV=1
CCG013139.1	2235	513	1823	9.86285184	28.51406647	1.531597034	Up	0	0	AAEL004520-PA OS=Aedes aegypti GN=AAEL004520 PE=4 SV=1
CCG015611.1	3381	288	1825	3.66024359	18.86981813	2.366068951	Up	0	0	AAEL009447-PA OS=Aedes aegypti GN=AAEL009447 PE=4 SV=1
CCG001103.1	984	284	1827	12.401834	64.90728828	2.387827003	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003570 PE=4 SV=1
CCG022490.1	24615	555	1827	0.96884837	2.594709391	1.421230161	Up	0	0	AAEL011664-PA OS=Aedes aegypti GN=AAEL011664 PE=4 SV=1
CCG016118.1	2535	3204	1827	54.3096764	25.19478172	-1.10808431	Down	1.94E-158	1.11E-157	AAEL000485-PA OS=Aedes aegypti GN=AAEL000485 PE=4 SV=1
CCG024588.1	2601	535	1828	8.83844983	24.56890809	1.474968478	Up	2.42E-14	8.43E-14	Uncharacterized protein OS=Anopheles darlingi GN=AND_11405 PE=3 SV=1
CCG004454.1	2524	691	1829	11.7639012	25.33228535	1.106610663	Up	2.46E-12	5.81E-12	AAEL002386-PA OS=Aedes aegypti GN=AAEL002386 PE=4 SV=1

CCG021323.2	8572	713	1831	3.57412748	7.467172745	1.062971013	Up	8.97E-14	2.84E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017756 PE=4 SV=1
CCG006079.1	435	338	1832	33.3879777	147.2265817	2.140637556	Up	1.44E-13	4.32E-13	AAEL005270-PA OS=Aedes aegypti GN=AAEL005270 PE=2 SV=1
CCG001481.1	1878	693	1834	15.8562439	34.13923301	1.106379585	Up	1.99E-12	4.75E-12	Syntaxin-5 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003088 PE=3 SV=1
CCG005546.1	2730	483	1835	7.60233766	23.49759629	1.627998173	Up	0	0	AAEL013334-PA OS=Aedes aegypti GN=AAEL013334 PE=4 SV=1
CCG005819.1	1667	663	1835	17.0899424	38.48136645	1.171012491	Up	1.08E-13	3.38E-13	Autophagy-related protein 3 OS=Aedes aegypti GN=AAEL000955 PE=3 SV=1
CCG003350.1	1393	277	1839	8.54459188	46.1509483	2.433276802	Up	0	0	AAEL005579-PA OS=Aedes aegypti GN=AAEL005579 PE=4 SV=1
CCG024460.2	4491	344	1839	3.29138025	14.31491227	2.120754214	Up	4.69E-13	1.22E-12	Flt-1 mRNA-splicing peccase BKRZ OS=Culex quinquefasciatus GN=CpipJ_CPIJ013139 PE=4 SV=1
CCG024437.1	2172	405	1840	8.01231239	29.61474644	1.886025157	Up	2.34E-13	6.64E-13	AAEL013022-PA OS=Aedes aegypti GN=AAEL013022 PE=4 SV=1
CCG023358.1	3699	743	1840	8.63111998	17.38935638	1.010584854	Up	0	0	AAEL000264-PA OS=Aedes aegypti GN=AAEL000264 PE=4 SV=1
CCG016327.2	1614	159	1843	4.23307794	39.91828011	3.237270604	Up	5.53E-14	1.84E-13	AAEL000262-PA OS=Aedes aegypti GN=AAEL000262 PE=4 SV=1
CCG025307.1	1611	424	1844	11.3092287	40.01431556	1.82301569	Up	0	0	S-hydroxyisobutyrate dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ000400 PE=4 SV=1
CCG001067.1	4809	437	1844	3.90471491	13.4046709	1.779446675	Up	5.07E-13	1.31E-12	AAEL012168-PA OS=Aedes aegypti GN=AAEL012168 PE=4 SV=1
CCG000852.1	2709	680	1844	10.7860537	23.79588866	1.141545208	Up	4.12E-13	1.09E-12	AAEL011545-PA (Fragment) OS=Aedes aegypti GN=AAEL011545 PE=4 SV=1
CCG026831.1	357	650	1845	78.2362115	180.6667245	1.207422397	Up	0	0	AAEL002263-PA OS=Aedes aegypti GN=AAEL002263 PE=4 SV=1

CCG007220.1	1881	239	1846	5.45973768	34.30780379	2.651633233	Up	6.72E-13	1.70E-12	AAEL005074-PA OS=Aedes aegypti GN=AAEL005074 PE=4 SV=1
CCG022469.1	246	286	1847	49.9566833	262.4712894	2.393410018	Up	0	0	AAEL006931-PA OS=Aedes aegypti GN=AAEL006931 PE=4 SV=1
CCG021919.1	4323	552	1847	5.48676695	14.9359096	1.444756898	Up	0	0	AAEL001742-PA OS=Aedes aegypti GN=AAEL001742 PE=4 SV=1
CCG013503.1	2366	729	1848	13.2396181	27.30468955	1.044287241	Up	4.15E-13	1.10E-12	AAEL012510-PA OS=Aedes aegypti GN=IKK2 PE=4 SV=1
CCG007303.1	1842	27	1849	0.62984953	35.09112582	5.799955211	Up	2.28E-12	5.40E-12	AAEL004948-PA OS=Aedes aegypti GN=AAEL004948 PE=3 SV=1
CCG026786.3	909	7327	1849	346.357806	71.10875	-2.284164187	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010605 PE=4 SV=1
CCG018184.1	1326	348	1850	11.2771249	48.77285975	2.112679263	Up	0	0	AAEL013248-PA OS=Aedes aegypti GN=AAEL013248 PE=3 SV=1
CCG022742.2	3077	514	1852	7.17791472	21.04086077	1.551557038	Up	1.57E-13	4.65E-13	Ubiquitin carboxyl-terminal hydrolase OS=Aedes aegypti GN=AAEL012219 PE=3 SV=1
CCG024673.1	3783	452	1853	5.13410522	17.12336422	1.737781407	Up	1.12E-13	3.49E-13	AAEL007312-PA OS=Aedes aegypti GN=AAEL007312 PE=4 SV=1
CCG021661.1	2761	482	1856	7.50141691	23.49966016	1.647404863	Up	0	0	AAEL012137-PA OS=Aedes aegypti GN=AAEL012137 PE=4 SV=1
CCG005047.1	738	522	1859	30.3932269	88.05885707	1.534718262	Up	0	0	AAEL013854-PA (Fragment) OS=Aedes aegypti GN=AAEL013854 PE=3 SV=1
CCG013617.1	378	409	1861	46.4937076	172.1093997	1.888218511	Up	3.15E-13	8.65E-13	Histone H2A OS=Anopheles gambiae GN=AGAP000306 PE=3 SV=3
CCG027188.1	12676	545	1861	1.84746808	5.132325108	1.474063124	Up	0	0	AAEL005388-PA OS=Aedes aegypti GN=AAEL005388 PE=4 SV=1
CCG028056.1	3141	426	1863	5.82779591	20.73456531	1.831015543	Up	4.37E-13	1.15E-12	AAEL001742-PA OS=Aedes aegypti GN=AAEL001742 PE=4 SV=1
CCG005326.1	3893	658	1865	7.26280128	16.74728646	1.205329345	Up	1.34E-12	3.25E-12	AAEL010520-PA OS=Aedes aegypti GN=AAEL010520 PE=4 SV=1

CCG015087.1	5764	3359	1865	25.040829	11.311101	-1.146542961	Down	2.43E-175	1.42E-174	AAEL007484-PA OS=Aedes aegypti GN=AAEL007484 PE=4 SV=1
CCG020927.1	2434	287	1866	5.06668604	26.80038803	2.403139548	Up	2.87E-13	7.99E-13	AAEL004083-PA OS=Aedes aegypti GN=AAEL004083 PE=4 SV=1
CCG005108.1	6381	626	1866	4.21549191	10.22287172	1.278027628	Up	0	0	AAEL006013-PA OS=Aedes aegypti GN=AAEL006013 PE=4 SV=1
CCG009900.1	1611	199	1867	5.30786914	40.51340952	2.932194796	Up	0	0	AAEL002978-PA OS=Aedes aegypti GN=AAEL002978 PE=4 SV=1
CCG005491.1	1737	52	1868	1.28637087	37.59473864	4.869152225	Up	0	0	AAEL013242-PA OS=Aedes aegypti GN=AAEL013242 PE=4 SV=1
CCG007732.3	2916	454	1868	6.69007528	22.39439678	1.743043456	Up	3.16E-13	8.66E-13	AAEL004791-PA OS=Aedes aegypti GN=AAEL004791 PE=4 SV=1
CCG010193.1	1629	314	1869	8.28268672	40.10866746	2.275743309	Up	4.97E-14	1.68E-13	AAEL001199-PA OS=Aedes aegypti GN=AAEL001199 PE=3 SV=1
CCG013616.1	1959	247	1870	5.41782769	33.37007533	2.622768527	Up	0	0	AAEL015216-PA OS=Aedes aegypti GN=AAEL015216 PE=3 SV=1
CCG017380.1	2205	85	1872	1.65642968	29.67886355	4.163286987	Up	0	0	AAEL002810-PA OS=Aedes aegypti GN=AAEL002810 PE=3 SV=1
CCG016733.1	1731	221	1872	5.4860262	37.80583138	2.784775364	Up	0	0	AAEL011061-PA OS=Aedes aegypti GN=AAEL011061 PE=4 SV=1
CCG015616.1	3060	317	1873	4.45143983	21.39766418	2.265109358	Up	2.22E-13	6.35E-13	AAEL007857-PA OS=Aedes aegypti GN=AAEL007857 PE=4 SV=1
CCG017608.1	3966	4694	1873	50.8572704	16.50954422	-1.623153738	Down	0	0	AAEL005384-PA OS=Aedes aegypti GN=AAEL005384 PE=4 SV=1
CCG016905.1	4635	604	1874	5.59950803	14.13415548	1.335813702	Up	2.31E-13	6.59E-13	AAEL005126-PA OS=Aedes aegypti GN=AAEL005126 PE=4 SV=1
CCG026531.1	6733	467	1876	2.98037518	9.740342674	1.708478577	Up	5.22E-14	1.75E-13	Transcription initiation factor TFIID subunit 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003658 PE=4 SV=1
CCG019437.1	2318	224	1877	4.1523816	28.30745708	2.769171217	Up	2.04E-13	5.87E-13	Purative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014589 PE=4 SV=1
CCG007362.1	1866	330	1878	7.59914921	35.18308884	2.210972337	Up	1.76E-13	5.16E-13	AAEL000575-PA OS=Aedes aegypti GN=AAEL000575 PE=3 SV=1

CCG027189.2	7601	246	1879	1.39067948	8.641836871	2.63554805	Up	2.51E-13	7.05E-13	AAEL014900-PA OS=Aedes aegypti GN=AAEL014900 PE=4 SV=1
CCG018805.1	3335	390	1880	5.02494648	19.70661479	1.971499837	Up	6.59E-13	1.67E-12	AAEL012388-PA OS=Aedes aegypti GN=AAEL012388 PE=4 SV=1
CCG010694.1	1533	560	1880	15.6967067	42.871207	1.449547133	Up	0	0	AAEL009113-PA OS=Aedes aegypti GN=AAEL009113 PE=4 SV=1
CCG010977.1	1734	625	1880	15.4879378	37.90170723	1.291117771	Up	0	0	-
CCG023355.1	1461	27	1881	0.79410187	45.00788406	5.824709835	Up	2.28E-12	5.41E-12	AAEL009330-PA OS=Aedes aegypti GN=AAEL009330 PE=4 SV=1
CCG025715.1	2277	686	1881	12.9456469	28.87857646	1.157532571	Up	2.38E-13	6.73E-13	AAEL014844-PA OS=Aedes aegypti GN=AAEL014844 PE=4 SV=1
CCG018767.1	2567	320	1883	5.35656996	25.64333275	2.259202394	Up	0	0	AAEL012830-PA OS=Aedes aegypti GN=AAEL012830 PE=2 SV=1
CCG005134.1	915	192	1884	9.01660005	71.97966496	2.996933953	Up	0	0	AAEL006317-PA OS=Aedes aegypti GN=AAEL006317 PE=4 SV=1
CCG013391.1	2052	86	1885	1.80087582	32.11323183	4.156397257	Up	0	0	AAEL010877-PA OS=Aedes aegypti GN=AAEL010877 PE=4 SV=1
CCG019811.1	2754	123	1889	1.91912758	23.97828062	3.643205685	Up	0	0	Zinc finger protein 467 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001514 PE=4 SV=1
CCG022123.1	2697	4091	1891	65.179527	24.51097567	-1.41099097	Down	1.28E-294	8.05E-294	AAEL003499-PA OS=Aedes aegypti GN=AAEL003499 PE=4 SV=1
CCG025758.1	961	110	1893	4.918492	68.86162116	3.807412186	Up	0	0	AAEL009773-PA OS=Aedes aegypti GN=AAEL009773 PE=4 SV=1
CCG009521.1	1824	114	1893	2.68560841	36.28071158	3.755881885	Up	8.48E-14	2.70E-13	AAEL009572-PA (Fragment) OS=Aedes aegypti GN=AAEL009572 PE=3 SV=1
CCG015294.1	3527	539	1893	6.56668187	18.76269292	1.514630437	Up	7.19E-13	1.81E-12	AAEL004790-PA OS=Aedes aegypti GN=AAEL004790 PE=4 SV=1
CCG006188.1	2922	151	1895	2.22054412	22.67143548	3.351890598	Up	0	0	AAEL014099-PA OS=Aedes aegypti GN=AAEL014099 PE=3 SV=1
CCG024508.1	1389	608	1895	18.8089263	47.69325737	1.342367823	Up	1.30E-12	3.16E-12	AAEL012325-PA OS=Aedes aegypti GN=AAEL012325 PE=4 SV=1

CCG022098.1	2700	683	1896	10.8697514	24.5484788	1.175314684	Up	1.46E-12	3.53E-12	CG1553, isoform B OS=Drosophila melanogaster GN=Nop17l PE=2 SV=1
CCG016286.1	1830	41	1897	0.9627099	36.23816997	5.234265163	Up	1.02E-13	3.20E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004372 PE=4 SV=1
CCG018983.1	1051	593	1897	24.2445791	63.09786017	1.379928873	Up	1.13E-12	2.78E-12	AAEL010582-PA OS=Aedes aegypti GN=GSTD11 PE=3 SV=1
CCG002413.1	1056	444	1898	18.0668202	62.83220579	1.798161614	Up	6.43E-13	1.64E-12	AAEL011069-PA OS=Aedes aegypti GN=AAEL011069 PE=4 SV=1
CCG023797.1	2161	464	1898	9.22626416	30.70375257	1.734596486	Up	0	0	AAEL015108-PA OS=Aedes aegypti GN=AAEL015108 PE=4 SV=1
CCG008904.2	1575	605	1899	16.5058346	42.14969371	1.352546062	Up	2.07E-13	5.93E-13	AAEL000819-PA OS=Aedes aegypti GN=AAEL000819 PE=4 SV=1
CCG014760.1	11971	634	1899	2.275734	5.545549043	1.284998364	Up	3.52E-13	9.51E-13	AAEL002645-PA OS=Aedes aegypti GN=AAEL002645 PE=4 SV=1
CCG003741.1	786	292	1901	15.9633111	84.54921647	2.405031462	Up	0	0	-
CCG014805.1	21488	346	1901	0.69189911	3.092688205	2.160227793	Up	2.24E-13	6.40E-13	AAEL013319-PA OS=Aedes aegypti GN=AAEL013319 PE=4 SV=1
CCG024379.1	3210	388	1901	5.19384954	20.70270534	1.994943178	Up	3.43E-13	9.27E-13	AAEL002290-PA OS=Aedes aegypti GN=AAEL002290 PE=4 SV=1
CCG009127.1	1815	4791	1903	113.425895	36.65322352	-1.629738048	Down	0	0	AAEL008672-PA OS=Aedes aegypti GN=AAEL008672 PE=3 SV=1
CCG021491.1	8431	123	1904	0.62688618	7.894740716	3.654616462	Up	0	0	AAEL010240-PA OS=Aedes aegypti GN=AAEL010240 PE=4 SV=1
CCG017200.1	2097	9405	1905	192.718338	31.75751896	-2.601323742	Down	0	0	Pyruvate kinase OS=Aedes aegypti GN=AAEL014913 PE=3 SV=1
CCG013199.1	1524	54	1908	1.52254965	43.76666147	4.845271158	Up	0	0	AAEL006121-PA OS=Aedes aegypti GN=AAEL006121 PE=3 SV=1
CCG006864.1	1536	5971	1908	167.039248	43.42473443	-1.943598194	Down	0	0	AAEL007881-PA OS=Aedes aegypti GN=AAEL007881 PE=4 SV=1
CCG020674.1	3669	318	1909	3.7242779	18.1889753	2.288031636	Up	1.20E-13	3.69E-13	Tetrapeptide repeat protein 37 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001407 PE=4 SV=1

CCG010663.1	2631	550	1909	8.98265072	25.36501344	1.497626783	Up	5.05E-13	1.31E-12	AAEL008134-PA OS=Aedes aegypti GN=AAEL008134 PE=3 SV=1
CCG017251.1	2967	543	1910	7.86402625	22.50431703	1.516861739	Up	1.00E-12	2.48E-12	AAEL010525-PA OS=Aedes aegypti GN=AAEL010525 PE=4 SV=1
CCG014850.1	31008	449	1912	0.62220752	2.155580018	1.792608377	Up	1.06E-12	2.60E-12	AAEL006523-PA OS=Aedes aegypti GN=AAEL006523 PE=4 SV=1
CCG011803.1	2883	513	1914	7.64601937	23.20851257	1.601873303	Up	0	0	Lon protease homolog OS=Aedes aegypti GN=AAEL006474 PE=3 SV=1
CCG027581.1	2139	247	1915	4.96190951	31.29738196	2.657074649	Up	0	0	AAEL015048-PA OS=Aedes aegypti GN=AAEL015048 PE=4 SV=1
CCG005176.1	2505	66	1917	1.13213672	26.75250163	4.562553706	Up	0	0	AAEL008706-PA OS=Aedes aegypti GN=AAEL008706 PE=4 SV=1
CCG020781.2	2085	175	1918	3.60657245	32.15826132	3.156489097	Up	3.55E-13	9.58E-13	AAEL004877-PA OS=Aedes aegypti GN=AAEL004877 PE=4 SV=1
CCG010943.1	1251	264	1918	9.06795359	53.5971022	2.563306089	Up	0	0	AAEL003013-PA OS=Aedes aegypti GN=AAEL003013 PE=3 SV=1
CCG027445.1	1857	528	1919	12.2175659	36.12543518	1.564058081	Up	0	0	AAEL001518-PA (Fragment) OS=Aedes aegypti GN=AAEL001518 PE=4 SV=1
CCG022971.1	2223	663	1920	12.8155349	30.19338345	1.236338739	Up	1.08E-13	3.38E-13	AAEL012941-PA OS=Aedes aegypti GN=AAEL012941 PE=4 SV=1
CCG011143.1	546	532	1922	41.8679465	123.0582563	1.555423389	Up	2.30E-13	6.57E-13	AAEL000950-PA OS=Aedes aegypti GN=AAEL000950 PE=4 SV=1
CCG023581.1	1162	390	1923	14.4218559	57.85263875	2.004125937	Up	6.59E-13	1.68E-12	Eukaryotic translation initiation factor 3 subunit G OS=Anopheles darlingi GN=AND_16989 PE=3 SV=1
CCG007605.1	1407	282	1924	8.6122709	47.80364215	2.472654935	Up	5.45E-13	1.40E-12	AAEL002166-PA OS=Aedes aegypti GN=AAEL002166 PE=4 SV=1
CCG010308.2	4188	653	1924	6.69991325	16.06010614	1.261267106	Up	5.96E-13	1.53E-12	AGAP004049-PA OS=Anopheles gambiae GN=AgaP_AGAP004049 PE=4 SV=5
CCG016097.1	906	309	1925	14.6552406	74.27669182	2.341492906	Up	6.25E-13	1.59E-12	AAEL006710-PA OS=Aedes aegypti GN=AAEL006710 PE=4 SV=1

CCG024721.2	4002	394	1925	4.23040366	16.81526307	1.990904115	Up	5.49E-13	1.41E-12	AAEL005303-PA OS=Aedes aegypti GN=AAEL005303 PE=4 SV=1
CCG015671.1	1531	7585	1925	212.884022	43.95472422	-2.275977531	Down	0	0	AAEL001429-PA OS=Aedes aegypti GN=AAEL001429 PE=4 SV=1
CCG013964.1	2189	166	1927	3.25855457	30.77414314	3.239414629	Up	1.83E-13	5.32E-13	AAEL003949-PA OS=Aedes aegypti GN=AAEL003949 PE=4 SV=1
CCG018226.1	1426	494	1927	14.8857285	47.24025199	1.666086829	Up	0	0	AAEL014413-PA OS=Aedes aegypti GN=CYP304C1 PE=3 SV=1
CCG010684.1	2074	198	1929	4.10222153	32.51423139	2.986594012	Up	0	0	AAEL011651-PA OS=Aedes aegypti GN=AAEL011651 PE=4 SV=1
CCG011472.1	2267	175	1930	3.31702848	29.76156779	3.165487224	Up	3.55E-13	9.57E-13	AAEL002104-PA OS=Aedes aegypti GN=AAEL002104 PE=4 SV=1
CCG018911.2	2688	411	1931	6.57014915	25.11325612	1.934451069	Up	0	0	AAEL003992-PA OS=Aedes aegypti GN=AAEL003992 PE=4 SV=1
CCG017012.1	1743	75	1932	1.84895588	38.74893329	4.389373892	Up	0	0	AAEL012618-PA OS=Aedes aegypti GN=AAEL012618 PE=4 SV=1
CCG007514.1	3531	48	1933	0.58412553	19.1374537	5.033976625	Up	5.35E-14	1.80E-13	Micröcephalin OS=Culex quinquefasciatus GN=CpipJ_CPIJ009302 PE=4 SV=1
CCG026258.1	2448	145	1934	2.54518444	27.61818108	3.439776193	Up	0	0	AAEL012546-PA OS=Aedes aegypti GN=AAEL012546 PE=3 SV=1
CCG006821.1	2160	310	1934	6.16695265	31.30060522	2.343560878	Up	2.14E-13	6.15E-13	Asparagine synthetase OS=Aedes aegypti GN=AAEL015631 PE=4 SV=1
CCG004781.1	4591	312	1934	2.92018236	14.72648819	2.334283064	Up	9.28E-14	2.93E-13	AAEL013391-PA OS=Aedes aegypti GN=AAEL013391 PE=4 SV=1
CCG022899.1	2928	577	1934	8.467738	23.09061041	1.447257775	Up	4.04E-13	1.07E-12	AAEL003229-PA OS=Aedes aegypti GN=AAEL003229 PE=4 SV=1
CCG023527.1	6660	6852	1935	44.2085017	10.15679663	-2.121878381	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_09210 PE=4 SV=1
CCG005366.1	1752	209	1939	5.12595578	38.68955403	2.91605116	Up	0	0	AAEL014307-PA OS=Aedes aegypti GN=AAEL014307 PE=4 SV=1
CCG020115.1	4831	386	1939	3.43330937	14.03106989	2.030953255	Up	2.52E-13	7.09E-13	AAEL015543-PA OS=Aedes aegypti GN=AAEL015543 PE=4 SV=1

CCG002838.1	3010	94	1941	1.34191198	22.5428622	4.070308755	Up	0	0	wvd-repeat protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011019 PE=4 SV=1
CCG009990.1	867	260	1941	12.8859642	78.26299333	2.602529794	Up	3.22E-13	8.81E-13	AAEL0008484-PA OS=Aedes aegypti GN=AAEL008484 PE=4 SV=1
CCG007783.1	2712	164	1942	2.59846478	25.03280733	3.268088685	Up	1.14E-13	3.54E-13	AAEL0008728-PA OS=Aedes aegypti GN=AAEL008728 PE=4 SV=1
CCG026809.1	3786	169	1942	1.91808905	17.93158307	3.224761253	Up	0	0	AAEL012637-PA OS=Aedes aegypti GN=AAEL012637 PE=4 SV=1
CCG003631.1	3957	433	1942	4.70202049	17.15667766	1.867417475	Up	9.43E-13	2.34E-12	AAEL004551-PA OS=Aedes aegypti GN=AAEL004551 PE=4 SV=1
CCG017390.1	15353	539	1942	1.50854471	4.42187022	1.551499226	Up	7.19E-13	1.81E-12	AAEL010378-PA OS=Aedes aegypti GN=AAEL010378 PE=4 SV=1
CCG019500.1	2076	157	1943	3.24963793	32.71865692	3.331762641	Up	0	0	AAEL002478-PA OS=Aedes aegypti GN=AAEL002478 PE=4 SV=1
CCG010777.1	4437	291	1943	2.8181638	15.30852643	2.441508046	Up	6.00E-14	1.99E-13	AAEL009871-PA OS=Aedes aegypti GN=AAEL009871 PE=4 SV=1
CCG024660.1	2805	685	1944	10.4935003	24.22776829	1.20716553	Up	4.65E-13	1.21E-12	AAEL009877-PA (Fragment) OS=Aedes aegypti GN=AAEL009877 PE=4 SV=1
CCG011007.1	1638	555	1945	14.5593423	41.51028591	1.511523682	Up	0	0	-
CCG023999.1	2352	597	1946	10.9068587	28.92381233	1.407022077	Up	0	0	AAEL009068-PA OS=Aedes aegypti GN=AAEL009068 PE=3 SV=1
CCG019201.1	2250	96	1947	1.83337534	30.25056217	4.044387872	Up	0	0	AAEL011269-PA OS=Aedes aegypti GN=AAEL011269 PE=4 SV=1
CCG019952.1	1084	780	1948	30.9191817	62.82170033	1.022760852	Up	1.45E-12	3.51E-12	AAEL012577-PA OS=Aedes aegypti GN=AAEL012577 PE=4 SV=1
CCG027743.1	1422	639	1949	19.3091845	47.91398131	1.311159458	Up	4.28E-13	1.13E-12	AAEL009837-PA OS=Aedes aegypti GN=AAEL009837 PE=4 SV=1
CCG006720.1	2306	6575	1949	122.517782	29.54626255	-2.0519436	Down	0	0	AAEL009414-PA OS=Aedes aegypti GN=AAEL009414 PE=4 SV=1
CCG024651.1	1422	3996	1950	120.750393	47.93856519	-1.332769255	Down	8.88E-264	5.50E-263	AAEL007703-PA OS=Aedes aegypti GN=AAEL007703 PE=4 SV=1

CCG028520.1	833	6260	1950	322.917814	81.83510169	-1.980375329	Down	0	0	NADH dehydrogenase OS=Ochlerotatus taeniorhynchus PE=2 SV=1
CCG024725.1	3002	379	1951	5.42489321	22.7193864	2.066257228	Up	1.40E-13	4.20E-13	AAEL014002-PA OS=Aedes aegypti GN=AAEL014002 PE=4 SV=1
CCG016278.1	2094	720	1951	14.7746938	32.5709637	1.14045817	Up	0	0	AAEL000077-PA OS=Aedes aegypti GN=AAEL000077 PE=4 SV=1
CCG022019.1	1251	11	1952	0.3778314	54.54720724	7.173618923	Up	2.67E-07	5.04E-07	AAEL011582-PA OS=Aedes aegypti GN=AAEL011582 PE=4 SV=1
CCG000753.2	2064	257	1952	5.35039816	33.06131602	2.627425992	Up	4.14E-13	1.09E-12	AAEL007672-PA OS=Aedes aegypti GN=AAEL007672 PE=4 SV=1
CCG026109.1	966	365	1952	16.2359763	70.64032739	2.121297888	Up	2.91E-13	8.05E-13	AAEL009037-PA (Fragment) OS=Aedes aegypti GN=AAEL009037 PE=4 SV=1
CCG001824.1	3743	291	1953	3.34068736	18.24031914	2.448914095	Up	6.00E-14	1.99E-13	AAEL001238-PA OS=Aedes aegypti GN=AAEL001238 PE=4 SV=1
CCG027936.1	1838	645	1953	15.0791506	37.14554654	1.300634087	Up	3.47E-13	9.37E-13	Peptide-O-fucosyltransferase OS=Glossina morsitans morsitans PE=2 SV=1
CCG014845.1	3091	421	1954	5.85255848	22.09915005	1.916851533	Up	4.57E-13	1.20E-12	AAEL010302-PA OS=Aedes aegypti GN=AAEL010302 PE=4 SV=1
CCG026601.2	1197	652	1955	23.4054026	57.09559824	1.286537942	Up	1.50E-12	3.61E-12	AAEL003844-PA OS=Aedes aegypti GN=GALE5 PE=4 SV=1
CCG010323.1	2223	172	1958	3.32469382	30.79096083	3.211213499	Up	1.35E-13	4.09E-13	AAEL014377-PA OS=Aedes aegypti GN=AAEL014377 PE=4 SV=1
CCG016771.1	1719	276	1960	6.8991546	39.8593499	2.530426686	Up	0	0	AAEL001879-PA OS=Aedes aegypti GN=AAEL001879 PE=4 SV=1
CCG023595.1	3752	333	1961	3.81367847	18.27110361	2.260308657	Up	4.41E-13	1.16E-12	AAEL003573-PA OS=Aedes aegypti GN=AAEL003573 PE=4 SV=1
CCG012727.1	784	11869	1961	650.520128	87.44028157	-2.895223744	Down	0	0	AAEL008871-PA OS=Aedes aegypti GN=AAEL008871 PE=2 SV=1
CCG007038.1	742	156	1962	9.03406819	92.43684505	3.355020311	Up	1.42E-13	4.26E-13	-
CCG024886.1	7800	536	1963	2.95279202	8.797832988	1.575068471	Up	0	0	AAEL006187-PA OS=Aedes aegypti GN=AAEL006187 PE=4 SV=1

CCG028407.1	9702	503	1964	2.22776505	7.076690948	1.667477828	Up	1.17E-13	3.61E-13	Transcription-associated protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ013348 PE=4 SV=1
CCG024534.2	3087	556	1967	7.73928489	22.27500175	1.525153373	Up	1.85E-13	5.35E-13	B-box type zinc-finger protein hcf1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001613 PE=4 SV=1
CCG003242.1	594	7174	1967	518.964438	115.7625091	-2.164467584	Down	0	0	AAEL010850-PA OS=Aedes aegypti GN=AAEL010850 PE=4 SV=1
CCG007201.1	1401	3555	1969	109.034551	49.13122553	-1.150073245	Down	2.29E-186	1.35E-185	AAEL001446-PA (Fragment) OS=Aedes aegypti GN=AAEL001446 PE=4 SV=1
CCG000955.1	3240	730	1970	9.68145255	21.25549544	1.134540464	Up	0	0	AAEL009068-PA OS=Aedes aegypti GN=AAEL009068 PE=3 SV=1
CCG013274.1	2482	196	1971	3.39325866	27.7609845	3.032315421	Up	7.93E-14	2.55E-13	AAEL001751-PA OS=Aedes aegypti GN=AAEL001751 PE=3 SV=1
CCG003872.1	3513	213	1972	2.60533831	19.62360427	2.91304742	Up	3.57E-13	9.60E-13	AAEL002437-PA OS=Aedes aegypti GN=AAEL002437 PE=4 SV=1
CCG014524.1	1611	436	1972	11.6293012	42.79188193	1.879572715	Up	1.61E-13	4.76E-13	AAEL010827-PA OS=Aedes aegypti GN=AAEL010827 PE=4 SV=1
CCG005704.1	1247	148	1974	5.09985623	55.3389241	3.439766113	Up	0	0	AAEL006037-PA OS=Aedes aegypti GN=AAEL006037 PE=4 SV=1
CCG010457.1	2266	340	1975	6.44735647	30.46893055	2.240559206	Up	8.86E-14	2.81E-13	AAEL006525-PA OS=Aedes aegypti GN=AAEL006525 PE=4 SV=1
CCG022269.1	2907	314	1978	4.64138172	23.78653989	2.357519166	Up	4.97E-14	1.68E-13	AAEL014494-PA OS=Aedes aegypti GN=AAEL014494 PE=3 SV=1
CCG006086.1	1116	5	1979	0.19251673	61.99142449	8.330941006	Up	1.59E-05	2.76E-05	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG023662.1	954	25	1979	1.12604126	72.51827016	6.009012911	Up	9.32E-12	2.08E-11	AAEL003647-PA (Fragment) OS=Aedes aegypti GN=GPRNPR3 PE=3 SV=2
CCG010493.1	5408	3233	1979	25.6880828	12.79260905	-1.005788691	Down	1.38E-136	7.73E-136	AAEL005627-PA OS=Aedes aegypti GN=AAEL005627 PE=4 SV=1
CCG001745.1	840	468	1980	23.9402807	82.40165239	1.783233199	Up	5.00E-13	1.30E-12	AAEL004109-PA OS=Aedes aegypti GN=AAEL004109 PE=4 SV=1

CCG026567.1	1343	159	1981	5.08725823	51.56541049	3.341443413	Up	5.53E-14	1.84E-13	AAEL006836-PA OS=Aedes aegypti GN=AAEL006836 PE=2 SV=1
CCG028338.1	2556	361	1981	6.06888662	27.09403219	2.158471342	Up	1.35E-13	4.08E-13	AAEL000730-PA OS=Aedes aegypti GN=AAEL000730 PE=4 SV=1
CCG018407.1	2590	124	1985	2.05723826	26.79234726	3.703040185	Up	1.61E-13	4.77E-13	AAEL004493-PA OS=Aedes aegypti GN=AAEL004493 PE=4 SV=1
CCG011384.1	3405	208	1985	2.62487659	20.37949468	2.956796778	Up	3.93E-13	1.05E-12	DNA-directed RNA polymerase OS=Anopheles darlingi GN=AND_15508 PE=3 SV=1
CCG014385.1	1749	8786	1986	215.855968	39.69533314	-2.443027673	Down	0	0	AAEL005269-PA OS=Aedes aegypti GN=AAEL005269 PE=3 SV=1
CCG014580.1	1941	117	1987	2.59013856	35.78675731	3.788324641	Up	9.44E-14	2.97E-13	AAEL005265-PA OS=Aedes aegypti GN=AAEL005265 PE=4 SV=1
CCG010452.1	2565	476	1987	7.97411059	27.08073916	1.763871598	Up	4.62E-13	1.21E-12	AAEL001598-PA OS=Aedes aegypti GN=AAEL001598 PE=4 SV=1
CCG011591.1	3718	334	1988	3.86011064	18.69205331	2.275710953	Up	3.15E-13	8.65E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003232 PE=4 SV=1
CCG017378.1	10452	14540	1988	59.7761138	6.649163244	-3.168324403	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007110 PE=4 SV=1
CCG021332.1	1203	542	1992	19.3595978	57.88602438	1.580166095	Up	0	0	AAEL013396-PA OS=Aedes aegypti GN=AAEL013396 PE=4 SV=1
CCG023340.1	1851	601	1993	13.9518155	37.64011108	1.431818018	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001780 PE=4 SV=1
CCG013675.1	1131	8593	1994	326.471202	61.63289468	-2.405183281	Down	0	0	AAEL004631-PA OS=Aedes aegypti GN=AAEL004631 PE=3 SV=1
CCG026145.1	649	2	1995	0.13241829	107.4603423	9.664486235	Up	9.36E-05	0.00015393	AAEL009504-PA OS=Aedes aegypti GN=AAEL009504 PE=4 SV=1
CCG007982.1	2589	741	1995	12.2984061	26.93772196	1.131156503	Up	0	0	AAEL004989-PA OS=Aedes aegypti GN=AAEL004989 PE=4 SV=1
CCG004180.1	969	3823	1995	169.528685	71.97292277	-1.23600325	Down	1.70E-224	1.03E-223	AAEL013305-PA OS=Aedes aegypti GN=AAEL013305 PE=3 SV=1

CCG001602.1	3379	360	1996	4.57801257	20.65010963	2.173356113	Up	5.08E-13	1.32E-12	AAEL011815-PA OS=Aedes aegypti GN=AAEL011815 PE=4 SV=1
CCG027431.1	2113	301	1998	6.12110275	33.05567297	2.433034395	Up	0	0	AAEL001129-PA OS=Aedes aegypti GN=AAEL001129 PE=4 SV=1
CCG011566.1	1134	268	1999	10.1551048	61.62398172	2.60128677	Up	0	0	AAEL009441-PA OS=Aedes aegypti GN=AAEL009441 PE=4 SV=1
CCG022518.1	2115	497	1999	10.0973797	33.04094339	1.710273919	Up	7.04E-14	2.29E-13	AAEL013817-PA (Fragment) OS=Aedes aegypti GN=AAEL013817 PE=4 SV=1
CCG026345.1	2181	537	1999	10.5798934	32.0410799	1.598597683	Up	3.67E-13	9.84E-13	AAEL004419-PA OS=Aedes aegypti GN=AAEL004419 PE=4 SV=1
CCG022802.1	11835	162386	1999	589.58034	5.904655282	-6.641691771	Down	0	0	ADP1,ATP carrier protein 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005941 PE=3 SV=1
CCG020757.1	3423	792	2000	9.94216471	20.42551959	1.038740868	Up	1.23E-12	3.00E-12	AAEL011927-PA OS=Aedes aegypti GN=AAEL011927 PE=4 SV=1
CCG006085.3	1983	1	2001	0.02166905	35.2755985	10.66881866	Up	0.000587	0.00091666	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG026289.1	2795	360	2002	5.53456331	25.03988197	2.177686366	Up	5.08E-13	1.32E-12	AAEL001729-PA OS=Aedes aegypti GN=AAEL001729 PE=4 SV=1
CCG003740.1	3588	540	2002	6.46701691	19.50570516	1.592723866	Up	9.26E-13	2.31E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008967 PE=4 SV=1
CCG019942.1	2838	203	2004	3.07359272	24.68512567	3.00564408	Up	9.28E-14	2.93E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ017587 PE=4 SV=1
CCG011201.1	2964	377	2005	5.4654487	23.64755227	2.113279012	Up	4.66E-13	1.21E-12	AAEL009428-PA (Fragment) OS=Aedes aegypti GN=AAEL009428 PE=4 SV=1
CCG005789.1	3708	170	2006	1.97002559	18.91216376	3.263028158	Up	3.53E-13	9.52E-13	AAEL003896-PA OS=Aedes aegypti GN=AAEL003896 PE=4 SV=1
CCG013476.1	2427	199	2007	3.52327037	28.90863679	3.036513485	Up	0	0	-
CCG004576.1	1470	161	2009	4.70620903	47.77631159	3.343658174	Up	0	0	AAEL010402-PA OS=Aedes aegypti GN=AAEL010402 PE=3 SV=1

CCG001271.1	3249	783	2009	10.355587	21.61624439	1.061706555	Up	0	0	AAEL000622-PA OS=Aedes aegypti GN=AAEL000622 PE=4 SV=1
CCG016851.1	1668	385	2010	9.91807424	42.125981	2.086578354	Up	7.93E-13	1.99E-12	Alkaline phosphatase OS=Aedes aegypti GN=AAEL003313 PE=3 SV=1
CCG020145.1	1401	7969	2010	244.415286	50.15427289	-2.284889992	Down	0	0	AAEL002175-PA OS=Aedes aegypti GN=AAEL002175 PE=4 SV=1
CCG006556.1	12684	397	2011	1.34492153	5.542501939	2.043015373	Up	8.02E-13	2.01E-12	AAEL000140-PA OS=Aedes aegypti GN=AAEL000140 PE=4 SV=1
CCG012056.1	1938	13178	2011	292.185326	36.27507461	-3.009833241	Down	0	0	AGAP004944-PA OS=Anopheles gambiae GN=AGAP004944 PE=3 SV=3
CCG026424.3	6315	818	2013	5.56599254	11.1434697	1.001487628	Up	0	0	AAEL010014-PA OS=Aedes aegypti GN=AAEL010014 PE=4 SV=1
CCG026576.1	2421	402	2014	7.1349993	29.0813587	2.027109481	Up	0	0	AAEL003146-PA OS=Aedes aegypti GN=AAEL003146 PE=4 SV=1
CCG011634.1	1758	9	2015	0.21998158	40.06878709	7.508952326	Up	1.09E-06	2.01E-06	AAEL004821-PA OS=Aedes aegypti GN=AAEL004821 PE=4 SV=1
CCG000855.1	5536	755	2015	5.86021489	12.72415601	1.118544493	Up	3.09E-13	8.51E-13	AAEL007689-PA OS=Aedes aegypti GN=AAEL007689 PE=4 SV=1
CCG027477.1	408	350	2016	36.8612919	172.7350146	2.228382015	Up	3.00E-14	1.04E-13	-
CCG012504.1	2278	535	2016	10.0916629	30.93761456	1.616198046	Up	2.42E-14	8.44E-14	AAEL008479-PA OS=Aedes aegypti GN=AAEL008479 PE=4 SV=1
CCG015278.1	1188	5	2018	0.18084905	59.38198866	8.359095568	Up	1.59E-05	2.76E-05	AAEL009875-PA OS=Aedes aegypti GN=AAEL009875 PE=4 SV=1
CCG013168.1	1119	249	2018	9.56162995	63.04361262	2.721021731	Up	0	0	AAEL000634-PA OS=Aedes aegypti GN=AAEL000634 PE=4 SV=1
CCG004859.1	1104	508	2018	19.7723054	63.90018345	1.692338976	Up	6.20E-13	1.58E-12	AAEL011849-PA OS=Aedes aegypti GN=AAEL011849 PE=4 SV=1
CCG015174.1	3165	674	2018	9.15058487	22.28935309	1.284418882	Up	0	0	AAEL003357-PA OS=Aedes aegypti GN=AAEL003357 PE=4 SV=1
CCG023674.1	3993	797	2018	8.5767289	17.66736853	1.042587749	Up	5.68E-13	1.46E-12	AAEL002061-PA OS=Aedes aegypti GN=AAEL002061 PE=3 SV=1
CCG008698.1	2194	424	2019	8.30408727	32.16990009	1.953817945	Up	0	0	AAEL009913-PA OS=Aedes aegypti GN=AAEL009913 PE=4 SV=1

CCG002063.1	11851	494	2019	1.79116099	5.955679757	1.733371168	Up	0	0	AAEL008796-PA OS=Aedes aegypti GN=AAEL008796 PE=4 SV=1
CCG008098.1	2421	183	2020	3.24802207	29.16799632	3.166752943	Up	3.05E-13	8.41E-13	AAEL011376-PA (Fragment) OS=Aedes aegypti GN=AAEL011376 PE=3 SV=1
CCG004874.1	1247	272	2020	9.37270875	56.62848362	2.59498994	Up	5.36E-13	1.38E-12	AAEL009730-PA OS=Aedes aegypti GN=AAEL009730 PE=4 SV=1
CCG008320.1	3843	422	2021	4.71850846	18.3842512	1.962067621	Up	5.52E-13	1.42E-12	AAEL013182-PA OS=Aedes aegypti GN=AAEL013182 PE=4 SV=1
CCG021193.1	1767	536	2022	13.034396	40.00318938	1.617791295	Up	0	0	AAEL010460-PA OS=Aedes aegypti GN=AAEL010460 PE=4 SV=1
CCG010862.1	1545	5424	2022	150.852971	45.75122047	-1.721260977	Down	0	0	26S proteasome regulatory chain 4 OS=Aedes aegypti GN=AAEL012095 PE=2 SV=1
CCG022993.1	1499	363	2023	10.4056129	47.17851495	2.18076807	Up	0	0	AAEL005070-PA OS=Aedes aegypti GN=AAEL005070 PE=4 SV=1
CCG028591.1	2097	315	2025	6.45468116	33.75799259	2.386811378	Up	4.28E-13	1.13E-12	AAEL000768-PA OS=Aedes aegypti GN=AAEL000768 PE=4 SV=1
CCG028490.1	1464	361	2027	10.5956791	48.40193102	2.19158855	Up	1.35E-13	4.08E-13	AAEL005476-PA OS=Aedes aegypti GN=AAEL005476 PE=4 SV=1
CCG014680.1	7266	376	2028	2.2235921	9.757140833	2.133566289	Up	3.71E-14	1.28E-13	AAEL000693-PA OS=Aedes aegypti GN=AAEL000693 PE=4 SV=1
CCG007315.1	1991	461	2030	9.94929566	35.64304462	1.840954275	Up	1.29E-13	3.95E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_11769 PE=4 SV=1
CCG006220.1	3417	141	2031	1.7731146	20.77853676	3.550736376	Up	0	0	AAEL011780-PA OS=Aedes aegypti GN=AAEL011780 PE=3 SV=1
CCG009512.1	4887	281	2031	2.47073776	14.52839372	2.555861408	Up	0	0	AAEL001384-PA (Fragment) OS=Aedes aegypti GN=AAEL001384 PE=4 SV=1
CCG025931.1	1748	352	2032	8.65294427	40.6379968	2.231566272	Up	5.16E-13	1.34E-12	ATP-dependent (S)-NAD(P)-hydrate dehydratase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005232 PE=3 SV=1

CCG006173.1	2336	79	2035	1.45317168	30.45380703	4.389345535	Up	0	0	AAEL011376-PA (Fragment) OS=Aedes aegypti GN=AAEL011376 PE=3 SV=1
CCG016201.1	2979	134	2036	1.93284473	23.89226301	3.627745859	Up	0	0	DNA topoisomerase OS=Aedes aegypti GN=AAEL007683 PE=3 SV=1
CCG008188.1	4320	519	2037	5.16233617	16.48379856	1.67495274	Up	0	0	AGAP006599-PA OS=Anopheles gambiae GN=AGAP006599 PE=4 SV=4
CCG019021.1	3800	653	2037	7.38400966	18.73947626	1.343604287	Up	5.96E-13	1.53E-12	-
CCG015441.2	1419	4144	2037	125.487372	50.18323452	-1.322264819	Down	3.79E-270	2.36E-269	AAEL009131-PA OS=Aedes aegypti GN=CYP6Z8 PE=3 SV=1
CCG004396.1	927	517	2038	23.9647819	76.85541323	1.68123107	Up	8.39E-14	2.69E-13	AAEL001606-PA (Fragment) OS=Aedes aegypti GN=GPRMTN PE=3 SV=2
CCG026052.1	6444	711	2038	4.74107407	11.05601615	1.22154579	Up	0	0	Plasma membrane calcium- transporting ATPase 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ005742 PE=3 SV=1
CCG000757.1	3078	278	2039	3.88095719	23.15787081	2.577018191	Up	2.11E-14	7.39E-14	AAEL008413-PA (Fragment) OS=Aedes aegypti GN=AAEL008413 PE=4 SV=1
CCG017400.1	4158	265	2040	2.73857135	17.1512469	2.646818091	Up	4.94E-13	1.28E-12	Serine/threonine-protein kinase 3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000288 PE=4 SV=1
CCG018847.1	4319	455	2040	4.52679538	16.51189734	1.866943906	Up	0	0	AAEL003519-PA OS=Aedes aegypti GN=AAEL003519 PE=4 SV=1
CCG024566.1	7857	762	2044	4.16735876	9.094402154	1.125845497	Up	5.84E-13	1.50E-12	AAEL009677-PA OS=Aedes aegypti GN=AAEL009677 PE=4 SV=1
CCG010994.1	4170	382	2047	3.93631622	17.16057375	2.124179763	Up	6.17E-13	1.58E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009355 PE=4 SV=1
CCG015290.1	1893	8067	2049	183.115081	37.83914903	-2.274799068	Down	0	0	AAEL004739-PA OS=Aedes aegypti GN=AAEL004739 PE=3 SV=1

CCG004805.1	1293	790	2050	26.2537435	55.42495544	1.078012555	Up	8.07E-13	2.02E-12	AAEL014187-PA OS=Aedes aegypti GN=AAEL014187 PE=4 SV=1
CCG016290.1	873	76	2051	3.74077873	82.1299263	4.456497467	Up	0	0	AAEL013721-PA OS=Aedes aegypti GN=AAEL013721 PE=4 SV=1
CCG014405.1	2779	608	2051	9.40107904	25.80044104	1.456497467	Up	1.30E-12	3.16E-12	AAEL005445-PA OS=Aedes aegypti GN=AAEL005445 PE=4 SV=1
CCG011802.2	2718	213	2052	3.36738538	26.3923414	2.970418599	Up	3.57E-13	9.60E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008798 PE=4 SV=1
CCG024078.1	792	703	2052	38.1410649	90.57371709	1.24774734	Up	2.48E-13	6.98E-13	AAEL006554-PA OS=Aedes aegypti GN=AAEL006554 PE=4 SV=1
CCG026014.2	14079	10170	2052	31.0392926	5.095133457	-2.606903839	Down	0	0	AAEL011129-PA OS=Aedes aegypti GN=AAEL011129 PE=3 SV=1
CCG001177.1	2817	588	2053	8.96918848	25.47722478	1.506158771	Up	8.31E-13	2.08E-12	AAEL009416-PA OS=Aedes aegypti GN=AAEL009416 PE=4 SV=1
CCG008926.1	1974	420	2056	9.14249673	36.4104443	1.993692235	Up	1.84E-13	5.32E-13	AAEL000569-PA OS=Aedes aegypti GN=AAEL000569 PE=4 SV=1
CCG011066.1	20902	683	2057	1.40409189	3.440301183	1.292897514	Up	1.46E-12	3.53E-12	AAEL003548-PA (Fragment) OS=Aedes aegypti GN=AAEL003548 PE=4 SV=1
CCG006361.2	5421	4005	2058	31.7457641	13.27137679	-1.258246057	Down	9.36E-242	5.74E-241	Kinesin-like protein Kif-1A OS=Culex quinquefasciatus GN=CpipJ_CPIJ004723 PE=3 SV=1
CCG002466.1	1782	696	2060	16.782792	40.41192489	1.26779833	Up	3.08E-13	8.48E-13	AAEL010065-PA OS=Aedes aegypti GN=AAEL010065 PE=2 SV=1
CCG027597.1	913	8013	2060	377.126488	78.87628713	-2.257384935	Down	0	0	AAEL013876-PA OS=Aedes aegypti GN=AAEL013876 PE=2 SV=1
CCG018681.1	2115	152	2062	3.08813223	34.08225376	3.464214308	Up	8.39E-14	2.69E-13	AAEL004842-PA OS=Aedes aegypti GN=AAEL004842 PE=4 SV=1
CCG000983.1	18455	403	2062	0.93832582	3.905931547	2.057505793	Up	0	0	Cation efflux protein/zinc transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ010101 PE=4 SV=1
CCG024489.1	28948	3584	2062	5.32000583	2.490119065	-1.095213101	Down	1.10E-173	6.38E-173	-

CCG002101.1	1074	240	2063	9.60217533	67.14983704	2.805950714	Up	0	0	AAEL006129-PA OS=Aedes aegypti GN=AAEL006129 PE=4 SV=1
CCG009304.2	12626	4297	2063	14.6238674	5.711937667	-1.356272752	Down	3.22E-291	2.02E-290	AAEL007778-PA OS=Aedes aegypti GN=AAEL007778 PE=4 SV=1
CCG007022.1	577	10466	2063	779.412899	124.9894714	-2.640581234	Down	0	0	AAEL007490-PA OS=Aedes aegypti GN=AAEL007490 PE=4 SV=1
CCG016160.1	7047	414	2064	2.52440331	10.23895037	2.020053502	Up	8.70E-13	2.17E-12	Activating signal cointegrator 1 complex subunit 3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006748 PE=4 SV=1
CCG012277.1	2400	464	2064	8.30748202	30.06411802	1.855559464	Up	0	0	AAEL011202-PB OS=Aedes aegypti GN=AAEL011202 PE=4 SV=1
CCG025548.1	29804	733	2064	1.05679826	2.420946291	1.195871071	Up	0	0	AGAP007375-PA OS=Anopheles gambiae GN=AGAP007375 PE=4 SV=4
CCG006022.1	8848	96	2065	0.46621774	8.158775038	4.129276769	Up	0	0	-
CCG009652.1	2730	111	2067	1.74712108	26.46840956	3.921220011	Up	0	0	AAEL013940-PA OS=Aedes aegypti GN=AAEL013940 PE=4 SV=1
CCG006944.1	3786	283	2067	3.21194794	19.08577868	2.570977634	Up	0	0	AAEL002925-PA OS=Aedes aegypti GN=AAEL002925 PE=4 SV=1
CCG026520.1	819	265	2068	13.9035161	88.27071595	2.666485125	Up	4.94E-13	1.28E-12	AAEL006743-PA OS=Aedes aegypti GN=AAEL006743 PE=2 SV=1
CCG003701.1	4212	780	2068	7.95735826	17.16375032	1.10900336	Up	1.45E-12	3.51E-12	AAEL005062-PA OS=Aedes aegypti GN=AAEL005062 PE=4 SV=1
CCG014397.1	2616	174	2069	2.85807868	27.64857593	3.274087638	Up	1.78E-13	5.20E-13	AAEL005827-PA (Fragment) OS=Aedes aegypti GN=AAEL005827 PE=4 SV=1
CCG007428.1	4875	6231	2070	54.9219316	14.84382214	-1.887519745	Down	0	0	AAEL004099-PA OS=Aedes aegypti GN=AAEL004099 PE=4 SV=1
CCG002483.1	1182	529	2071	19.2309557	61.25092318	1.67130113	Up	0	0	AAEL001017-PA OS=Aedes aegypti GN=AAEL001017 PE=4 SV=1
CCG012061.1	43842	5473	2071	5.36411107	1.651352383	-1.699691099	Down	0	0	SLC4-like anion exchanger OS=Aedes aegypti PE=2 SV=1
CCG013709.1	3747	202	2073	2.31648956	19.34040773	3.061606122	Up	0	0	AAEL000685-PA OS=Aedes aegypti GN=AAEL000685 PE=4 SV=1

CCG024804.1	2615	330	2073	5.42256689	27.71262247	2.353495391	Up	1.76E-13	5.16E-13	AAEL013205-PA OS=Aedes aegypti GN=AAEL013205 PE=4 SV=1
CCG004215.1	840	4170	2073	213.31404	86.27203303	-1.306014063	Down	1.13E-266	7.01E-266	AAEL005238-PA OS=Aedes aegypti GN=AAEL005238 PE=4 SV=1
CCG010543.1	1172	577	2074	21.1548949	61.86302562	1.548085874	Up	4.04E-13	1.07E-12	AAEL001623-PA OS=Aedes aegypti GN=AAEL001623 PE=2 SV=1
CCG017439.1	1526	248	2075	6.98328583	47.53500937	2.767012515	Up	5.86E-14	1.95E-13	AAEL006235-PA OS=Aedes aegypti GN=AAEL006235 PE=4 SV=1
CCG012726.1	1686	432	2075	11.0100388	43.02397645	1.966321323	Up	7.71E-13	1.94E-12	AAEL008864-PA OS=Aedes aegypti GN=AAEL008864 PE=4 SV=1
CCG016700.1	1150	735	2077	27.4632652	63.1376877	1.200998265	Up	2.17E-13	6.21E-13	AAEL009217-PA OS=Aedes aegypti GN=AAEL009217 PE=4 SV=1
CCG017923.1	1797	179	2079	4.28023511	40.4442167	3.240171469	Up	0	0	AAEL009572-PA (Fragment) OS=Aedes aegypti GN=AAEL009572 PE=3 SV=1
CCG024072.1	3288	612	2080	7.9980163	22.11472497	1.467293174	Up	6.59E-13	1.68E-12	AAEL012045-PA (Fragment) OS=Aedes aegypti GN=AAEL012045 PE=4 SV=1
CCG012447.2	2379	813	2080	14.6844869	30.56461357	1.057569475	Up	0	0	Probable membrane protein OS=Aedes aegypti PE=2 SV=1
CCG006391.1	2082	221	2081	4.56114858	34.94148605	2.937471894	Up	0	0	Acyl-coenzyme A oxidase OS=Aedes aegypti GN=AAEL000370 PE=3 SV=1
CCG002446.2	1785	696	2081	16.7545856	40.75527953	1.282430958	Up	3.08E-13	8.48E-13	AAEL011055-PA OS=Aedes aegypti GN=AAEL011055 PE=3 SV=1
CCG016161.1	4364	652	2083	6.41985952	16.68608857	1.378032174	Up	1.50E-12	3.61E-12	AAEL014059-PA OS=Aedes aegypti GN=AAEL014059 PE=4 SV=1
CCG022820.1	4848	5512	2084	48.8550283	15.02744406	-1.700907407	Down	0	0	AAEL003855-PA OS=Aedes aegypti GN=AAEL003855 PE=4 SV=1
CCG006581.1	2016	835	2086	17.7974843	36.17210583	1.023204259	Up	0	0	AAEL012683-PA OS=Aedes aegypti GN=AAEL012683 PE=4 SV=1
CCG015407.1	3852	351	2087	3.91546647	18.94027093	2.274200869	Up	4.44E-15	1.60E-14	Cell division cycle and apoptosis regulator protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ018512 PE=4 SV=1

CCG001092.1	2259	463	2088	8.80698855	32.31203271	1.875350817	Up	6.16E-13	1.57E-12	AAEL001009-PA (Fragment) OS=Aedes aegypti GN=AAEL001009 PE=4 SV=1
CCG020474.1	4248	482	2089	4.87556782	17.191111115	1.818020645	Up	0	0	AAEL001397-PA OS=Aedes aegypti GN=AAEL001397 PE=3 SV=1
CCG010782.1	4935	593	2089	5.16333387	14.79794127	1.519021686	Up	1.13E-12	2.78E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002903 PE=4 SV=1
CCG002846.1	664	191	2091	12.36027	110.086983	3.154861722	Up	6.68E-14	2.20E-13	AAEL003469-PA OS=Aedes aegypti GN=AAEL003469 PE=4 SV=1
CCG016191.1	5410	499	2091	3.96338218	13.51160014	1.769394545	Up	0	0	AAEL011534-PA OS=Aedes aegypti GN=AAEL011534 PE=4 SV=1
CCG010209.1	48867	834	2091	0.73335295	1.495851121	1.028386977	Up	4.58E-13	1.20E-12	AAEL001731-PA OS=Aedes aegypti GN=AAEL001731 PE=4 SV=1
CCG011957.1	1833	670	2092	15.7063405	39.89782597	1.344963055	Up	8.60E-13	2.15E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_03313 PE=4 SV=1
CCG005140.1	2355	681	2092	12.425643	31.05423143	1.321469352	Up	1.31E-14	4.67E-14	Transporter (Fragment) OS=Aedes aegypti GN=AAEL010883 PE=3 SV=1
CCG013877.1	4023	320	2093	3.4179257	18.18734111	2.411741705	Up	0	0	AAEL005480-PA OS=Aedes aegypti GN=AAEL005480 PE=4 SV=1
CCG008767.3	3351	553	2093	7.09109616	21.83457872	1.62253413	Up	7.29E-13	1.83E-12	AAEL010433-PA (Fragment) OS=Aedes aegypti GN=AAEL010433 PE=4 SV=1
CCG000704.1	2421	282	2094	5.00514877	30.23652687	2.594807578	Up	5.45E-13	1.40E-12	AAEL003146-PA OS=Aedes aegypti GN=AAEL003146 PE=4 SV=1
CCG024139.1	2792	703	2094	10.8193852	26.21870758	1.276978052	Up	2.48E-13	6.98E-13	Inhibitor of kappa kinase epsilon OS=Culex quinquefasciatus GN=CpipJ_CPIJ006852 PE=4 SV=1
CCG018734.1	7791	126	2095	0.69492832	9.400281073	3.757767809	Up	1.34E-13	4.06E-13	AAEL010069-PA OS=Aedes aegypti GN=AAEL010069 PE=4 SV=1
CCG028156.1	5667	236	2096	1.7894578	12.9296891	2.853093156	Up	4.60E-13	1.20E-12	AAEL011505-PA (Fragment) OS=Aedes aegypti GN=AAEL011505 PE=4 SV=1

CCG026469.1	3208	354	2096	4.74167271	22.84056986	2.268130655	Up	0	0	AAEL003839-PA OS=Aedes aegypti GN=AAEL003839 PE=4 SV=1
CCG017160.1	7584	3539	2097	20.0514097	9.666074155	-1.052701697	Down	7.15E-161	4.11E-160	AAEL005017-PA OS=Aedes aegypti GN=AAEL005017 PE=4 SV=1
CCG012746.1	23282	3997	2099	7.37694482	3.151680395	-1.226902202	Down	6.63E-232	4.04E-231	AAEL004972-PA OS=Aedes aegypti GN=AAEL004972 PE=4 SV=1
CCG022057.1	2013	185	2101	3.94903175	36.48650745	3.20779219	Up	0	0	Zinc finger protein 250 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001374 PE=4 SV=1
CCG014220.1	2850	745	2101	11.2324394	25.77099632	1.198077035	Up	0	0	AAEL012594-PA (Fragment) OS=Aedes aegypti GN=AAEL012594 PE=4 SV=1
CCG020864.1	11462	823	2101	3.08533341	6.407899101	1.05442503	Up	7.96E-13	1.99E-12	Axotactin OS=Culex quinquefasciatus GN=CpipJ_CPIJ006837 PE=4 SV=1
CCG018615.3	582	9019	2101	665.883224	126.1981778	-2.399578115	Down	0	0	Acyl carrier protein OS=Aedes aegypti GN=AAEL010611 PE=3 SV=1
CCG006801.1	1288	668	2102	22.2855456	57.05147343	1.356155865	Up	9.28E-13	2.31E-12	AAEL012250-PA OS=Aedes aegypti GN=AAEL012250 PE=4 SV=1
CCG002590.1	5197	4196	2102	34.6932858	14.13936844	-1.294938805	Down	8.48E-265	5.26E-264	AAEL003267-PA (Fragment) OS=Aedes aegypti GN=AAEL003267 PE=4 SV=2
CCG013857.1	984	42	2103	1.83407404	74.71265859	5.348228916	Up	7.66E-14	2.47E-13	AAEL006813-PA OS=Aedes aegypti GN=AAEL006813 PE=4 SV=1
CCG024070.1	2529	460	2103	7.81576826	29.06969397	1.895056288	Up	0	0	AAEL010912-PA OS=Aedes aegypti GN=AAEL010912 PE=4 SV=1
CCG026293.1	2871	664	2103	9.93796718	25.60684641	1.365506907	Up	4.96E-13	1.29E-12	AAEL006876-PA OS=Aedes aegypti GN=AAEL006876 PE=4 SV=1
CCG016956.1	2876	680	2104	10.1597425	25.57448342	1.331841257	Up	4.12E-13	1.09E-12	Sodium/hydrogen exchanger OS=Aedes aegypti GN=NHE8 PE=2 SV=1
CCG013985.1	3088	224	2106	3.11697557	23.84136363	2.935248003	Up	2.04E-13	5.88E-13	AAEL004228-PA OS=Aedes aegypti GN=AAEL004228 PE=4 SV=1

CCG010491.1	1039	259	2108	10.711416	70.92593593	2.727164068	Up	0	0	AAEL005638-PA OS=Aedes aegypti GN=AAEL005638 PE=4 SV=1
CCG002318.4	8288	302	2108	1.56574081	8.89141499	2.505567616	Up	4.59E-13	1.20E-12	AAEL003075-PA OS=Aedes aegypti GN=AAEL003075 PE=4 SV=1
CCG020802.1	1887	295	2109	6.71757907	39.07101522	2.540085439	Up	2.32E-13	6.62E-13	AAEL012002-PA OS=Aedes aegypti GN=AAEL012002 PE=4 SV=1
CCG028136.1	1881	744	2109	16.9960035	39.19564365	1.205497773	Up	0	0	AAEL012151-PA (Fragment) OS=Aedes aegypti GN=AAEL012151 PE=4 SV=1
CCG005765.1	1173	542	2110	19.8547282	62.88317476	1.663191446	Up	0	0	AAEL009363-PA OS=Aedes aegypti GN=AAEL009363 PE=4 SV=1
CCG025521.1	2034	4627	2110	97.7487522	36.26448574	-1.430520896	Down	0	0	Acyl-coenzyme A oxidase OS=Aedes aegypti GN=AAEL008574 PE=3 SV=1
CCG003904.1	2578	120	2113	2.00014281	28.65276913	3.84049966	Up	0	0	AAEL000999-PA OS=Aedes aegypti GN=AAEL000999 PE=3 SV=1
CCG023381.1	2232	216	2115	4.15836141	33.12578646	2.99386765	Up	1.38E-13	4.17E-13	AAEL011118-PA (Fragment) OS=Aedes aegypti GN=AAEL011118 PE=3 SV=1
CCG026583.1	1629	489	2116	12.8988338	45.40927787	1.815746461	Up	1.91E-13	5.52E-13	AAEL000433-PA OS=Aedes aegypti GN=AAEL000433 PE=3 SV=1
CCG013034.1	2522	359	2118	6.11662757	29.35829905	2.262960044	Up	0	0	AAEL000750-PA OS=Aedes aegypti GN=AAEL000750 PE=4 SV=1
CCG016018.1	1308	627	2119	20.5978774	56.63347743	1.459159443	Up	3.40E-13	9.23E-13	AAEL008744-PA OS=Aedes aegypti GN=AAEL008744 PE=4 SV=1
CCG017413.1	2875	462	2121	6.90504953	25.79008871	1.901093068	Up	0	0	AAEL006395-PA OS=Aedes aegypti GN=AAEL006395 PE=4 SV=1
CCG022428.1	2115	142	2122	2.88496563	35.07397792	3.603775025	Up	2.62E-13	7.34E-13	DNA polymerase OS=Aedes aegypti GN=AAEL009851 PE=3 SV=1
CCG016973.1	3921	153	2122	1.67670732	18.9190164	3.496134302	Up	0	0	AAEL011215-PA OS=Aedes aegypti GN=AAEL011215 PE=4 SV=1
CCG016903.1	1743	310	2124	7.64235096	42.59975896	2.478756849	Up	2.14E-13	6.15E-13	AAEL005110-PA OS=Aedes aegypti GN=AAEL005110 PE=4 SV=1
CCG024480.1	2815	620	2125	9.46402681	26.38946293	1.479435924	Up	4.74E-13	1.23E-12	AAEL010837-PA OS=Aedes aegypti GN=AAEL010837 PE=4 SV=1

CCG005910.1	1323	258	2133	8.37958543	56.36130337	2.749754199	Up	1.69E-13	4.97E-13	AAEL005626-PA OS=Aedes aegypti GN=AAEL005628 PE=4 SV=1
CCG020620.1	1101	22	2134	0.85861413	67.75745925	6.302226046	Up	8.38E-11	1.80E-10	AAEL004978-PA OS=Aedes aegypti GN=AAEL004978 PE=2 SV=1
CCG000634.1	2197	145	2135	2.83596337	33.9717437	3.582424468	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ016156 PE=4 SV=1
CCG001321.1	2393	597	2135	10.7199881	31.18926908	1.540746437	Up	0	0	AAEL010549-PA OS=Aedes aegypti GN=AAEL010549 PE=4 SV=1
CCG009209.1	2229	90	2136	1.73498255	33.49972148	4.271156039	Up	0	0	AAEL002569-PA OS=Aedes aegypti GN=AAEL002569 PE=4 SV=1
CCG016400.1	2853	264	2137	3.97616892	26.18501138	2.719290277	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009380 PE=4 SV=1
CCG020804.1	1746	351	2138	8.63824562	42.80687041	2.309032121	Up	4.44E-15	1.60E-14	AAEL001171-PA OS=Aedes aegypti GN=AAEL001171 PE=4 SV=1
CCG019509.1	1668	90	2139	2.31851086	44.82958874	4.273180875	Up	0	0	AAEL006024-PA OS=Aedes aegypti GN=AAEL006024 PE=4 SV=1
CCG020708.1	4296	297	2139	2.97067299	17.40590177	2.55071485	Up	2.44E-13	6.86E-13	AAEL005432-PA OS=Aedes aegypti GN=AAEL005432 PE=4 SV=1
CCG009538.1	3795	601	2140	6.80495665	19.71296767	1.534487104	Up	0	0	AAEL006499-PA OS=Aedes aegypti GN=AAEL006499 PE=4 SV=1
CCG023464.1	33949	521	2141	0.65943715	2.204650227	1.741242722	Up	0	0	AAEL003883-PA (Fragment) OS=Aedes aegypti GN=AAEL003883 PE=4 SV=1
CCG010882.2	1743	702	2141	17.306227	42.94071748	1.311055064	Up	6.78E-13	1.71E-12	AAEL003416-PA OS=Aedes aegypti GN=AAEL003416 PE=4 SV=1
CCG021494.1	804	11126	2142	594.628442	93.13511051	-2.674591422	Down	0	0	AAEL010230-PA OS=Aedes aegypti GN=AAEL010230 PE=4 SV=1
CCG010814.1	3879	69	2143	0.76434949	19.31311862	4.659204882	Up	0	0	AAEL009238-PA OS=Aedes aegypti GN=AAEL009238 PE=4 SV=1
CCG012752.1	3431	325	2143	4.07028964	21.83491318	2.423433431	Up	1.19E-13	3.66E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003370 PE=4 SV=1

CCG009029.1	1365	185	2144	5.82373693	54.90882447	3.237020934	Up	0	0	ATP-dependent (S)-NAD(P) ⁺ -hydrate dehydratase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005232 PE=3 SV=1
CCG005285.1	3207	325	2146	4.35458801	23.39272278	2.425451657	Up	1.19E-13	3.66E-13	AAEL010526-PA OS=Aedes aegypti GN=AAEL010526 PE=4 SV=1
CCG020653.1	3461	260	2148	3.22800665	21.69615097	2.748723669	Up	3.22E-13	8.81E-13	AAEL010081-PA OS=Aedes aegypti GN=AAEL010081 PE=4 SV=1
CCG011136.1	2555	294	2149	4.94446261	29.40326293	2.572090626	Up	0	0	AAEL014701-PA OS=Aedes aegypti GN=AAEL014701 PE=4 SV=1
CCG003852.1	1493	8682	2151	249.874907	50.36520652	-2.310706694	Down	0	0	AAEL006833-PA OS=Aedes aegypti GN=AAEL006833 PE=2 SV=1
CCG005104.1	642	344	2157	23.0242815	117.4532757	2.35085891	Up	4.69E-13	1.22E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003833 PE=4 SV=1
CCG008944.1	999	185	2161	7.95735826	75.62045656	3.2484151	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003010 PE=4 SV=1
CCG011238.1	2952	710	2163	10.3348616	25.61475361	1.309455939	Up	9.64E-14	3.03E-13	Ubiquitin carboxyl-terminal hydrolase OS=Culex quinquefasciatus GN=CpipJ_CPIJ010867 PE=3 SV=1
CCG003643.1	1872	389	2164	8.92907413	40.41117037	2.178171643	Up	0	0	AGAP011732-PA OS=Anopheles gambiae GN=AGAP011732 PE=4 SV=3
CCG007390.1	2394	826	2164	14.8258149	31.59971217	1.091800016	Up	1.49E-13	4.45E-13	AGAP005404-PA OS=Anopheles gambiae GN=AGAP005404 PE=4 SV=3
CCG026795.1	2914	326	2165	4.80718376	25.97277598	2.433736359	Up	6.17E-13	1.58E-12	AAEL003815-PA OS=Aedes aegypti GN=AAEL003815 PE=4 SV=1
CCG008525.1	2616	171	2167	2.80880146	28.95817499	3.365944127	Up	3.42E-13	9.25E-13	AAEL011827-PA OS=Aedes aegypti GN=AAEL011827 PE=4 SV=1
CCG025458.1	8509	746	2167	3.76723728	8.902877631	1.240764822	Up	0	0	Cholinephosphate cytidyl transferase isoform B2 OS=Aedes aegypti GN=CCT PE=2 SV=1

CCG018557.1	2079	448	2169	9.25947143	36.47162209	1.977772619	Up	0	0	AAEL002240-PA OS=Aedes aegypti GN=AAEL002240 PE=4 SV=1
CCG020061.1	1500	158	2173	4.52614538	50.64289028	3.484004915	Up	0	0	AAEL014146-PA OS=Aedes aegypti GN=AAEL014146 PE=4 SV=1
CCG027101.1	2572	503	2173	8.40349009	29.53512264	1.813371073	Up	1.17E-13	3.61E-13	AAEL011204-PA OS=Aedes aegypti GN=AAEL011204 PE=4 SV=1
CCG028418.1	2861	258	2174	3.87493587	26.56389154	2.777222173	Up	1.69E-13	4.97E-13	AAEL002910-PA OS=Aedes aegypti GN=AAEL002910 PE=4 SV=1
CCG027765.1	894	672	2174	32.2993978	85.01039564	1.396132006	Up	0	0	AAEL005862-PA (Fragment) OS=Aedes aegypti GN=AAEL005862 PE=4 SV=1
CCG020651.1	2906	111	2175	1.64130783	26.16457398	3.994697023	Up	0	0	AAEL010086-PA OS=Aedes aegypti GN=AAEL010086 PE=3 SV=1
CCG009302.1	3061	449	2175	6.30297643	24.83967722	1.978541255	Up	1.06E-12	2.60E-12	AAEL007417-PA OS=Aedes aegypti GN=AAEL007417 PE=4 SV=1
CCG006348.1	2301	158	2176	2.95055109	33.05919611	3.485995297	Up	0	0	AAEL014161-PA OS=Aedes aegypti GN=AAEL014161 PE=4 SV=1
CCG014986.1	2595	646	2176	10.6968973	29.31376118	1.45438569	Up	1.08E-12	2.64E-12	AAEL006950-PA OS=Aedes aegypti GN=AAEL006950 PE=4 SV=1
CCG009246.1	1341	141	2178	4.51807053	56.77787234	3.65155009	Up	0	0	AAEL012427-PA OS=Aedes aegypti GN=AAEL012427 PE=4 SV=1
CCG024199.1	957	740	2181	33.2263361	79.66980318	1.261705798	Up	0	0	AAEL001800-PA OS=Aedes aegypti GN=AAEL001800 PE=4 SV=1
CCG017562.1	3117	91	2183	1.25449017	24.48313064	4.286614979	Up	0	0	AAEL013725-PA OS=Aedes aegypti GN=AAEL013725 PE=4 SV=1
CCG025021.1	2840	727	2187	10.9996469	26.92033497	1.291239155	Up	1.67E-12	4.00E-12	AAEL014095-PA OS=Aedes aegypti GN=AAEL014095 PE=4 SV=1
CCG021045.1	2718	338	2189	5.34355051	28.15440318	2.397490007	Up	1.44E-13	4.32E-13	DNA ligase (Fragment) OS=Anopheles gambiae GN=AGAP009222 PE=3 SV=3
CCG026527.2	1944	423	2190	9.34989596	39.38200933	2.074514505	Up	1.18E-13	3.65E-13	AAEL009674-PA OS=Aedes aegypti GN=AAEL009674 PE=4 SV=1
CCG028579.1	3284	283	2191	3.70293389	23.32325956	2.65502873	Up	0	0	AAEL011382-PA OS=Aedes aegypti GN=AAEL011382 PE=4 SV=1

CCG021071.1	1530	5292	2192	148.624729	50.08401483	-1.56925206	Down	0	0	Catalase OS=Aedes aegypti GN=CAT1B PE=3 SV=1
CCG003166.2	1667	272	2193	7.01125844	45.9889028	2.713540459	Up	5.36E-13	1.38E-12	AAEL010747-PA OS=Aedes aegypti GN=AAEL010747 PE=4 SV=1
CCG014148.1	1627	626	2194	16.5329157	47.14103211	1.511642167	Up	0	0	AAEL010473-PA OS=Aedes aegypti GN=AAEL010473 PE=4 SV=1
CCG001831.1	7698	70	2196	0.39073544	9.972509196	4.673692526	Up	0	0	AAEL010069-PA OS=Aedes aegypti GN=AAEL010069 PE=4 SV=1
CCG020089.1	3024	7747	2197	110.081526	25.39792793	-2.115789666	Down	0	0	Curagen IV alpha 1 chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ005294 PE=4 SV=1
CCG006812.1	1487	374	2198	10.8074517	51.67336405	2.257394415	Up	1.37E-13	4.12E-13	AAEL001048-PA OS=Aedes aegypti GN=AAEL001048 PE=3 SV=1
CCG024414.1	870	486	2199	24.0037828	88.36005819	1.880132589	Up	0	0	AAEL005536-PA OS=Aedes aegypti GN=AAEL005536 PE=4 SV=1
CCG006570.1	2674	682	2199	10.9593714	28.74841085	1.391317164	Up	1.08E-12	2.66E-12	AAEL005728-PA OS=Aedes aegypti GN=AAEL005728 PE=3 SV=1
CCG019133.1	1710	4028	2199	101.217597	44.95511732	-1.170902875	Down	5.13E-217	3.09E-216	AAEL002296-PA OS=Aedes aegypti GN=AAEL002296 PE=3 SV=1
CCG027878.1	4571	527	2202	4.95406916	16.84054374	1.765252806	Up	8.05E-13	2.01E-12	TBC1 domain family OS=Culex quinquefasciatus GN=CpipJ_CPIJ004857 PE=4 SV=1
CCG026413.3	1591	854	2203	23.0648355	48.40545803	1.069474724	Up	1.02E-12	2.51E-12	Proteasome subunit alpha type OS=Aedes aegypti GN=AAEL009898 PE=3 SV=1
CCG024798.1	1164	20408	2206	753.373148	66.25254172	-3.507316904	Down	0	0	AAEL010814-PA OS=Aedes aegypti GN=AAEL010814 PE=3 SV=1
CCG018239.1	1680	75	2208	1.91829172	45.94516376	4.58201897	Up	0	0	Rap35 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007568 PE=4 SV=1
CCG025665.1	6318	829	2208	5.63816239	12.21713756	1.115609369	Up	8.95E-14	2.83E-13	AAEL012079-PA (Fragment) OS=Aedes aegypti GN=AAEL012079 PE=4 SV=1
CCG004187.1	2719	360	2211	5.6892624	28.42690325	2.320943417	Up	5.08E-13	1.32E-12	AAEL012519-PA OS=Aedes aegypti GN=AAEL012519 PE=4 SV=1

CCG001243.1	1316	428	2211	13.9749593	58.73309266	2.071329527	Up	1.52E-13	4.51E-13	AAEL005579-PA OS=Aedes aegypti GN=AAEL005579 PE=4 SV=1
CCG021932.1	2673	380	2212	6.10867907	28.92918377	2.243593266	Up	0	0	AAEL014012-PA OS=Aedes aegypti GN=AAEL014012 PE=4 SV=1
CCG020956.1	1512	8961	2213	254.663883	51.16578472	-2.315343083	Down	0	0	AAEL008167-PA OS=Aedes aegypti GN=AAEL008167 PE=3 SV=1
CCG027945.1	1816	767	2214	18.1485608	42.61983743	1.231669943	Up	0	0	AGAP006127-PA OS=Anopheles gambiae GN=AGAP006127 PE=4 SV=1
CCG005210.1	935	1	2216	0.04595694	82.85298538	10.81605537	Up	0.000587	0.00091797	AAEL017520-PA OS=Aedes aegypti GN=FOHSDR-2 PE=2 SV=1
CCG008068.1	2148	492	2216	9.84222972	36.06496337	1.873540865	Up	0	0	AGAP005928-PA OS=Anopheles gambiae GN=AGAP005928 PE=4 SV=1
CCG023814.2	1305	516	2216	16.9903318	59.36210063	1.804828114	Up	9.75E-14	3.07E-13	AAEL012117-PA (Fragment) OS=Aedes aegypti GN=AAEL012117 PE=3 SV=1
CCG012912.1	2988	4606	2216	66.2378171	25.92621865	-1.353243326	Down	0	0	AAEL005290-PA OS=Aedes aegypti GN=AAEL005290 PE=4 SV=1
CCG013266.1	3727	450	2217	5.18818905	20.79487513	2.002925067	Up	9.36E-13	2.33E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009380 PE=4 SV=1
CCG007599.1	2280	223	2218	4.20274159	34.00765697	3.016456954	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006088 PE=4 SV=1
CCG016723.1	4065	631	2218	6.67008673	19.07440538	1.515860659	Up	0	0	Sit-1000 mo GTPase activating protein 1,3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014404 PE=4 SV=1
CCG010623.2	1713	469	2221	11.7646267	45.32535476	1.945862769	Up	0	0	AAEL001968-PA OS=Aedes aegypti GN=AAEL001968 PE=4 SV=1
CCG001399.1	906	502	2222	23.8088375	85.73652427	1.848412751	Up	6.14E-13	1.57E-12	SKCAZ and CDKNTA-interacting protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012086 PE=4 SV=1

CCG001279.1	1558	611	2222	16.8514171	49.85705455	1.564927735	Up	2.21E-13	6.34E-13	AAEL007001-PA OS=Aedes aegypti GN=AAEL007001 PE=4 SV=1
CCG022836.1	2199	288	2223	5.62768693	35.33981322	2.650680435	Up	0	0	AAEL012115-PA OS=Aedes aegypti GN=AAEL012115 PE=4 SV=1
CCG010479.1	3872	753	2223	8.35645924	20.07031231	1.264099382	Up	6.29E-13	1.60E-12	AAEL008198-PA (Fragment) OS=Aedes aegypti GN=AAEL008198 PE=4 SV=1
CCG013610.1	3084	380	2226	5.29458468	25.23253051	2.252695473	Up	0	0	AAEL010204-PA OS=Aedes aegypti GN=AAEL010204 PE=4 SV=1
CCG005920.1	2001	302	2227	6.48518733	38.90658789	2.584794307	Up	4.59E-13	1.20E-12	CTCF-like protein OS=Aedes aegypti PE=2 SV=1
CCG010710.1	1866	314	2228	7.23070561	41.74010753	2.529225972	Up	4.97E-14	1.68E-13	Transporter OS=Aedes aegypti GN=AAEL003624 PE=3 SV=1
CCG016382.1	3141	438	2231	5.99195917	24.83028191	2.051000942	Up	4.14E-13	1.09E-12	AAEL014315-PA OS=Aedes aegypti GN=AAEL014315 PE=4 SV=1
CCG018776.1	3984	593	2231	6.39584654	19.57628401	1.613899708	Up	1.13E-12	2.78E-12	AAEL000737-PB (Fragment) OS=Aedes aegypti GN=AAEL000737 PE=4 SV=1
CCG011403.1	5480	763	2232	5.98282984	14.23848061	1.250895269	Up	4.81E-13	1.25E-12	AAEL003275-PA OS=Aedes aegypti GN=AAEL003275 PE=4 SV=2
CCG017360.1	3714	285	2234	3.29735443	21.02767644	2.672908565	Up	4.57E-13	1.20E-12	F-box and leucine-rich repeat protein 11 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008371 PE=4 SV=1
CCG021446.1	2572	255	2235	4.26021863	30.37781827	2.834018883	Up	0	0	AAEL003539-PA OS=Aedes aegypti GN=AAEL003539 PE=4 SV=1
CCG009391.1	690	170	2236	10.5867462	113.2850824	3.419626741	Up	3.53E-13	9.52E-13	AAEL008578-PA OS=Aedes aegypti GN=AAEL008578 PE=4 SV=1
CCG028517.1	1673	4	2237	0.10273696	46.74337426	8.829662745	Up	2.95E-05	5.03E-05	AAEL006123-PA OS=Aedes aegypti GN=AAEL006123 PE=3 SV=1
CCG012675.1	1366	6	2237	0.18873968	57.24865676	8.244700244	Up	8.38E-06	1.48E-05	AAEL001401-PA OS=Aedes aegypti GN=AAEL001401 PE=4 SV=1
CCG023494.1	453	100	2238	9.48559263	172.707778	4.186451335	Up	0	0	Putative salivary secreted peptide OS=Aedes albopictus PE=2 SV=1

CCG012705.1	1765	482	2238	11.7345111	44.32669882	1.917418189	Up	0	0	AAEL001626-PA OS=Aedes aegypti GN=AAEL001626 PE=4 SV=1
CCG010555.2	4831	229	2239	2.0368597	16.20194198	2.991748229	Up	2.46E-13	6.93E-13	AAEL009067-PA OS=Aedes aegypti GN=AAEL009067 PE=4 SV=1
CCG003175.1	2112	646	2239	13.1432048	37.060408	1.495561662	Up	1.08E-12	2.64E-12	AAEL010723-PA (Fragment) OS=Aedes aegypti GN=AAEL010723 PE=4 SV=1
CCG001539.1	1804	132	2240	3.14412692	43.40717293	3.787202101	Up	1.46E-13	4.36E-13	AAEL002142-PA OS=Aedes aegypti GN=AAEL002142 PE=4 SV=1
CCG023700.1	1274	154	2240	5.19414374	61.46510202	3.56480968	Up	1.82E-13	5.28E-13	AAEL003403-PA (Fragment) OS=Aedes aegypti GN=AAEL003403 PE=4 SV=1
CCG003828.1	1890	356	2241	8.09377012	41.45052817	2.356506706	Up	7.93E-13	1.99E-12	AAEL009882-PA OS=Aedes aegypti GN=AAEL009882 PE=4 SV=1
CCG009526.1	2129	628	2241	12.6749616	36.79732186	1.537619388	Up	1.62E-13	4.78E-13	AAEL011442-PA OS=Aedes aegypti GN=AAEL011442 PE=4 SV=1
CCG023144.1	1392	401	2243	12.378494	56.33003937	2.186068682	Up	1.05E-13	3.28E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_23192 PE=4 SV=1
CCG022787.1	2502	87	2245	1.49415144	31.36743859	4.391869438	Up	0	0	AAEL005281-PA OS=Aedes aegypti GN=AAEL005281 PE=4 SV=1
CCG002506.1	1803	640	2245	15.2527067	43.52819265	1.512884839	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007543 PE=4 SV=1
CCG007594.1	3410	807	2246	10.1690838	23.02530488	1.179030553	Up	9.50E-13	2.36E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005947 PE=4 SV=1
CCG003245.1	1874	72	2247	1.6509183	41.91635427	4.666172612	Up	0	0	AGAP013200-PA OS=Anopheles gambiae GN=AgaP_AGAP013200 PE=4 SV=1
CCG028274.1	2353	513	2247	9.36824218	33.38344577	1.833282597	Up	0	0	AAEL006363-PA OS=Aedes aegypti GN=AAEL006363 PE=4 SV=1
CCG017904.2	1890	555	2248	12.6180967	41.58000327	1.720395563	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004720 PE=4 SV=1

CCG006965.1	2951	624	2248	9.08611128	26.63036468	1.551337305	Up	0	0	AAEL012743-PA OS=Aedes aegypti GN=AAEL012743 PE=4 SV=1
CCG017993.1	1739	769	2248	19.0015675	45.19045784	1.249899736	Up	6.41E-13	1.63E-12	NMD protein OS=Aedes aegypti PE=2 SV=1
CCG018761.1	4404	702	2249	6.84939911	17.85221718	1.382053929	Up	6.78E-13	1.71E-12	AAEL010022-PA OS=Aedes aegypti GN=AAEL010022 PE=4 SV=1
CCG020507.1	926	252	2252	11.6937075	85.01732105	2.862024392	Up	5.42E-14	1.81E-13	-
CCG025250.1	2310	557	2252	10.3611005	34.08053649	1.717770799	Up	5.29E-13	1.37E-12	AAEL001848-PA (Fragment) OS=Aedes aegypti GN=AAEL001848 PE=4 SV=1
CCG014048.1	3846	876	2254	9.78717824	20.48776803	1.065797944	Up	1.44E-12	3.49E-12	AAEL011480-PA OS=Aedes aegypti GN=AAEL011480 PE=4 SV=1
CCG012612.1	2628	752	2256	12.2957536	30.0098449	1.287275705	Up	1.67E-12	4.00E-12	AAEL000198-PA OS=Aedes aegypti GN=AAEL000198 PE=4 SV=1
CCG007643.1	1047	10	2259	0.41040816	75.42573756	7.521853664	Up	5.43E-07	1.01E-06	AAEL005606-PA OS=Aedes aegypti GN=AAEL005606 PE=4 SV=1
CCG015940.1	5799	128	2259	0.94846112	13.617994	3.843781759	Up	4.15E-14	1.42E-13	AAEL011505-PA (Fragment) OS=Aedes aegypti GN=AAEL011505 PE=4 SV=1
CCG026368.1	2636	280	2260	4.56431172	29.97181544	2.715137244	Up	0	0	AAEL014423-PA (Fragment) OS=Aedes aegypti GN=AAEL014423 PE=4 SV=1
CCG026633.2	1642	4752	2260	124.355773	48.11553319	-1.36989886	Down	0	0	AAEL003453-PA OS=Aedes aegypti GN=AAEL003453 PE=4 SV=1
CCG004104.1	786	468	2261	25.5850328	100.5606409	1.974693761	Up	5.00E-13	1.30E-12	60S ribosomal protein L2/L8 OS=Aedes aegypti GN=AAEL000987 PE=2 SV=1
CCG001501.1	2191	12155	2262	238.382987	36.09111002	-2.723565855	Down	0	0	AAEL010037-PA OS=Aedes aegypti GN=AAEL010037 PE=3 SV=1
CCG000413.3	2718	506	2266	7.99951645	29.14475908	1.865251775	Up	3.38E-13	9.20E-13	AAEL003911-PA OS=Aedes aegypti GN=AAEL003911 PE=4 SV=1
CCG004792.1	3213	870	2266	11.6351289	24.65467014	1.083373759	Up	0	0	AAEL000622-PA OS=Aedes aegypti GN=AAEL000622 PE=4 SV=1
CCG025253.1	1241	451	2267	15.6159148	63.86012364	2.031898256	Up	1.51E-12	3.65E-12	AAEL007469-PA OS=Aedes aegypti GN=AAEL007469 PE=4 SV=1

CCG004051.1	2964	528	2267	7.65452762	26.73765636	1.80448776	Up	0	0	AAEL009428-PA (Fragment) OS=Aedes aegypti GN=AAEL009428 PE=4 SV=1
CCG013303.1	2706	759	2267	12.0524865	29.28692293	1.280925804	Up	1.54E-12	3.70E-12	AGAP007911-PA (Fragment) OS=Anopheles gambiae GN=AGAP007911 PE=4 SV=3
CCG002316.1	3954	467	2269	5.07507994	20.06078149	1.982875358	Up	5.22E-14	1.75E-13	AAEL003079-PA (Fragment) OS=Aedes aegypti GN=AAEL003079 PE=3 SV=1
CCG014559.1	2805	759	2269	11.6271047	28.27819251	1.282198022	Up	1.54E-12	3.70E-12	AAEL013138-PA OS=Aedes aegypti GN=AAEL013138 PE=4 SV=1
CCG026012.1	1872	318	2273	7.29934594	42.44666832	2.539812217	Up	1.20E-13	3.69E-13	AAEL011114-PA OS=Aedes aegypti GN=AAEL011114 PE=4 SV=1
CCG006080.1	1506	199	2275	5.6779397	52.80881783	3.217339413	Up	0	0	Ubiquitin carboxyl-terminal hydrolase OS=Aedes aegypti GN=AAEL002698 PE=3 SV=1
CCG020337.1	1121	163	2276	6.2480524	70.97684026	3.505869892	Up	1.80E-13	5.25E-13	AAEL015143-PA OS=Aedes aegypti GN=AAEL015143 PE=2 SV=1
CCG019113.1	969	98	2277	4.34575231	82.14653892	4.240521936	Up	0	0	AAEL002142-PA OS=Aedes aegypti GN=AAEL002142 PE=4 SV=1
CCG028477.1	2467	122	2278	2.12497269	32.28007884	3.925127898	Up	8.04E-14	2.58E-13	AAEL010342-PA OS=Aedes aegypti GN=AAEL010342 PE=4 SV=1
CCG007593.1	2398	271	2279	4.8560459	33.22348322	2.774349372	Up	0	0	AAEL012489-PA OS=Aedes aegypti GN=AAEL012489 PE=4 SV=1
CCG006420.1	1434	1	2280	0.02996495	55.58219738	10.85713131	Up	0.000587	0.00091586	Tubulin alpha-2 chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ004534 PE=3 SV=1
CCG005913.1	3054	638	2281	8.97665052	26.10996376	1.540351321	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010636 PE=4 SV=1
CCG001803.4	2759	920	2281	14.3284363	28.90171414	1.012273884	Up	1.78E-12	4.26E-12	AAEL006744-PA (Fragment) OS=Aedes aegypti GN=AAEL006744 PE=4 SV=1
CCG024462.1	972	275	2283	12.1570751	82.10879205	2.75574054	Up	4.31E-13	1.14E-12	AAEL009153-PA OS=Aedes aegypti GN=AAEL009153 PE=4 SV=1

CCG013046.1	2871	327	2285	4.8941495	27.82294059	2.507144828	Up	6.36E-13	1.62E-12	AAEL011240-PA OS=Aedes aegypti GN=AAEL011240 PE=4 SV=1
CCG011874.1	2103	3808	2287	77.8072988	38.0169182	-1.033263909	Down	9.11E-168	5.28E-167	AAEL003588-PA OS=Aedes aegypti GN=AAEL003588 PE=4 SV=1
CCG016462.1	951	239	2289	10.7989133	84.1424769	2.961948143	Up	6.72E-13	1.70E-12	AAEL004305-PA OS=Aedes aegypti GN=AAEL004305 PE=4 SV=1
CCG006297.1	2004	163	2290	3.49504328	39.94733224	3.514716933	Up	1.80E-13	5.24E-13	AAEL013741-PA OS=Aedes aegypti GN=AAEL013741 PE=4 SV=1
CCG027501.1	2007	346	2290	7.40783666	39.88762023	2.428816859	Up	2.24E-13	6.40E-13	AAEL010781-PA OS=Aedes aegypti GN=AAEL010781 PE=4 SV=1
CCG020466.1	14498	395	2291	1.17071632	5.524169684	2.238366104	Up	0	0	AAEL014249-PA OS=Aedes aegypti GN=AAEL014249 PE=4 SV=1
CCG020242.1	1020	358	2293	15.0815343	78.5875771	2.381518066	Up	1.30E-13	3.97E-13	AAEL007055-PA OS=Aedes aegypti GN=AAEL007055 PE=3 SV=1
CCG017526.1	2925	624	2294	9.16687672	27.41685023	1.580560661	Up	0	0	AAEL010018-PA (Fragment) OS=Aedes aegypti GN=AAEL010018 PE=4 SV=1
CCG024663.1	1482	15	2296	0.43491634	54.15938156	6.960329535	Up	1.48E-08	2.92E-08	AAEL004396-PA (Fragment) OS=Aedes aegypti GN=GPROAR4 PE=3 SV=2
CCG017524.1	4716	444	2296	4.04549664	17.0195512	2.072804264	Up	6.43E-13	1.64E-12	AAEL014851-PA (Fragment) OS=Aedes aegypti GN=AAEL014851 PE=4 SV=1
CCG024912.1	3553	607	2296	7.34101574	22.59054418	1.621667424	Up	0	0	AAEL011480-PA OS=Aedes aegypti GN=AAEL011480 PE=4 SV=1
CCG001292.1	1488	722	2297	20.8495621	53.96449042	1.371993318	Up	0	0	AAEL007022-PA OS=Aedes aegypti GN=AAEL007022 PE=4 SV=1
CCG004446.1	2371	882	2298	15.9845238	33.88195699	1.083841441	Up	1.00E-12	2.48E-12	AAEL001824-PA OS=Aedes aegypti GN=AAEL001824 PE=4 SV=1
CCG005006.1	393	878	2299	95.998542	204.5014715	1.091026825	Up	8.17E-13	2.04E-12	AAEL015100-PA OS=Aedes aegypti GN=AAEL015100 PE=3 SV=1
CCG004348.1	7841	5841	2299	32.0094656	10.24985057	-1.642895715	Down	0	0	AAEL005017-PA OS=Aedes aegypti GN=AAEL005017 PE=4 SV=1
CCG009672.1	23302	8207	2299	15.1340062	3.449020612	-2.133535283	Down	0	0	Glutathione peroxidase OS=Aedes aegypti GN=GPx PE=2 SV=1

CCG013500.1	3792	218	2302	2.47030647	21.22203405	3.102800997	Up	0	0	AAEL012508-PA OS=Aedes aegypti GN=AAEL012508 PE=4 SV=1
CCG015864.1	2283	375	2302	7.05810358	35.24921293	2.320238537	Up	0	0	AAEL007638-PA OS=Aedes aegypti GN=AAEL007638 PE=4 SV=1
CCG023410.1	4181	552	2302	5.67311493	19.24753722	1.762460865	Up	0	0	Translocase of outer membrane 34 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001395 PE=4 SV=1
CCG016639.1	1431	7	2305	0.21019437	56.3094535	8.065509317	Up	4.31E-06	7.70E-06	AAEL005325-PA OS=Aedes aegypti GN=AAEL005325 PE=4 SV=1
CCG001891.1	1737	237	2305	5.86288262	46.3896534	2.98412099	Up	2.32E-13	6.61E-13	AAEL011180-PA OS=Aedes aegypti GN=AAEL011180 PE=4 SV=1
CCG004440.1	2043	220	2310	4.6271863	39.52697961	3.094630627	Up	4.61E-13	1.20E-12	Queuine tRNA-ribosyltransferase OS=Aedes aegypti GN=AAEL010993 PE=3 SV=1
CCG026235.1	2721	214	2311	3.37946461	29.69076723	3.135147762	Up	0	0	B-box type zinc-finger protein hci-1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001613 PE=4 SV=1
CCG014040.1	2382	239	2311	4.31140494	33.91627944	2.97574794	Up	6.72E-13	1.70E-12	AAEL002082-PA (Fragment) OS=Aedes aegypti GN=AAEL002082 PE=4 SV=1
CCG021789.1	2979	192	2316	2.76944916	27.17803592	3.294770241	Up	0	0	AAEL006610-PA OS=Aedes aegypti GN=AAEL006610 PE=4 SV=1
CCG010348.1	2064	87	2318	1.81122428	39.26031277	4.438034559	Up	0	0	Putative uncharacterized protein GLEAN_08099 OS=Tribolium castaneum GN=GLEAN_08099 PE=4 SV=1
CCG002136.1	4425	172	2318	1.67023601	18.31260691	3.4547133	Up	1.35E-13	4.09E-13	AAEL014694-PA OS=Aedes aegypti GN=AAEL014694 PE=4 SV=1
CCG023472.1	1325	4868	2318	157.869183	61.15719665	-1.368135398	Down	0	0	Acetyl-CoA acetyltransferase OS=Aedes aegypti PE=2 SV=1
CCG013000.1	2155	352	2321	7.01872231	37.65111851	2.423412393	Up	5.16E-13	1.34E-12	Tubulin gamma chain OS=Aedes aegypti GN=AAEL012843 PE=3 SV=1
CCG027886.2	1635	490	2321	12.8777798	49.62578617	1.946206072	Up	0	0	AAEL001830-PA OS=Aedes aegypti GN=AAEL001830 PE=4 SV=1

CCG018404.1	1613	559	2322	14.8915571	50.32431411	1.756760988	Up	3.87E-13	1.03E-12	AGAP009439-PA (Fragment) OS=Anopheles gambiae GN=AGAP009439 PE=4 SV=4
CCG019866.1	3726	862	2322	9.94093163	21.78559277	1.131921402	Up	0	0	AAEL003433-PA OS=Aedes aegypti GN=AAEL003433 PE=3 SV=1
CCG010939.1	1829	3788	2325	88.993633	44.43848742	-1.001892411	Down	1.13E-158	6.48E-158	AAEL012845-PA OS=Aedes aegypti GN=AAEL012845 PE=4 SV=1
CCG020133.1	2087	168	2326	3.45899157	38.96164436	3.493631163	Up	1.45E-13	4.33E-13	AGAP001938-PA OS=Anopheles gambiae GN=AgaP_AGAP001938 PE=4 SV=4
CCG025014.1	1287	934	2326	31.183941	63.18022671	1.018669846	Up	3.46E-13	9.35E-13	AAEL009470-PA (Fragment) OS=Aedes aegypti GN=AAEL009470 PE=4 SV=1
CCG011710.1	2172	440	2327	8.70473445	37.45299726	2.105208986	Up	0	0	AAEL010723-PA (Fragment) OS=Aedes aegypti GN=AAEL010723 PE=4 SV=1
CCG004553.1	1281	469	2327	15.7320886	63.50344266	2.013124587	Up	0	0	AAEL001888-PA OS=Aedes aegypti GN=AAEL001888 PE=4 SV=1
CCG010226.1	4908	784	2329	6.86395109	16.58879923	1.273098284	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001658 PE=4 SV=1
CCG016734.1	2067	138	2330	2.86880666	39.40628199	3.779902987	Up	1.49E-13	4.45E-13	AAEL011062-PA OS=Aedes aegypti GN=AAEL011062 PE=4 SV=1
CCG022841.1	19137	923	2330	2.0724808	4.256298525	1.038240606	Up	0	0	AAEL012151-PA (Fragment) OS=Aedes aegypti GN=AAEL012151 PE=4 SV=1
CCG012298.1	4224	570	2331	5.79847271	19.29160586	1.734228384	Up	7.69E-13	1.93E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_20180 PE=4 SV=1
CCG027243.1	3165	502	2332	6.81542078	25.75756759	1.918121723	Up	6.14E-13	1.57E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013215 PE=4 SV=1
CCG017706.1	2214	803	2332	15.5847773	36.82145503	1.2404091	Up	2.24E-13	6.41E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_11718 PE=4 SV=1

CCG005472.1	1716	285	2333	7.1365818	47.52777372	2.735465687	Up	4.57E-13	1.20E-12	AAEL004489-PA OS=Aedes aegypti GN=AAEL004489 PE=4 SV=1
CCG007011.1	3402	273	2336	3.44818858	24.00427235	2.799380622	Up	1.53E-14	5.41E-14	AAEL003896-PA OS=Aedes aegypti GN=AAEL003896 PE=4 SV=1
CCG026382.1	4449	565	2336	5.45693415	18.35525613	1.750030705	Up	4.06E-13	1.08E-12	AAEL002260-PA (Fragment) OS=Aedes aegypti GN=AAEL002260 PE=4 SV=1
CCG000005.1	7405	4605	2336	26.7218944	11.02802627	-1.276847678	Down	4.20E-284	2.63E-283	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002312 PE=4 SV=1
CCG000673.1	1365	188	2337	5.9181759	59.85164309	3.338166371	Up	1.80E-13	5.24E-13	AAEL003645-PA OS=Aedes aegypti GN=AAEL003645 PE=3 SV=1
CCG015968.1	531	244	2337	19.7450381	153.8559187	2.962017885	Up	2.78E-13	7.73E-13	Histone H2A OS=Anopheles gambiae GN=AGAP000306 PE=3 SV=3
CCG001255.2	1345	25834	2337	825.338382	60.74163035	-3.764228197	Down	0	0	AAEL013613-PA OS=Aedes aegypti GN=AAEL013613 PE=4 SV=1
CCG008287.1	1668	842	2339	21.6909572	49.02122864	1.176312927	Up	0	0	AAEL013132-PB OS=Aedes aegypti GN=AAEL013132 PE=4 SV=1
CCG019096.1	603	8369	2339	596.37431	135.6010106	-2.136850182	Down	0	0	AAEL002309-PA OS=Aedes aegypti GN=TPX4 PE=4 SV=1
CCG025204.3	1611	14134	2339	376.992073	50.75568552	-2.892892843	Down	0	0	AAEL014616-PA OS=Aedes aegypti GN=CYP9J27 PE=3 SV=1
CCG001876.1	2907	586	2341	8.66194169	28.1518149	1.700465569	Up	0	0	AAEL014709-PA OS=Aedes aegypti GN=AAEL014709 PE=3 SV=1
CCG011805.1	1465	628	2341	18.4197907	55.86165592	1.600601674	Up	1.62E-13	4.77E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013237 PE=4 SV=1
CCG012693.1	1161	377	2342	13.9531352	70.51876331	2.337417851	Up	4.66E-13	1.21E-12	AAEL009569-PA OS=Aedes aegypti GN=AAEL009569 PE=4 SV=1
CCG013546.1	1675	15515	2344	398.015184	48.92071687	-3.024306018	Down	0	0	AAEL009951-PA OS=Aedes aegypti GN=AAEL009951 PE=4 SV=1
CCG002338.1	2520	844	2347	14.3914508	32.55836333	1.177816142	Up	1.09E-12	2.67E-12	AAEL005998-PB (Fragment) OS=Aedes aegypti GN=AAEL005998 PE=4 SV=1

CCG010155.2	10850	607	2349	2.40392893	7.568386372	1.654591496	Up	0	0	AAEL008687-PA OS=Aedes aegypti GN=AAEL008687 PE=4 SV=1
CCG016772.1	2272	410	2352	7.75422147	36.18920201	2.222505449	Up	1.25E-12	3.04E-12	CK31 complex subunit b OS=Culex quinquefasciatus GN=CpipJ_CPIJ000780 PE=4 SV=1
CCG006311.1	1680	802	2353	20.5129328	48.96239598	1.255140383	Up	7.06E-13	1.78E-12	Apyrase OS=Culex quinquefasciatus GN=CpipJ_CPIJ011013 PE=3 SV=1
CCG012803.2	3072	888	2353	12.4209389	26.7763103	1.108182943	Up	0	0	AAEL002782-PA (Fragment) OS=Aedes aegypti GN=AAEL002782 PE=4 SV=1
CCG018570.1	3461	431	2354	5.35104178	23.77688053	2.15166775	Up	0	0	AGAP006599-PA OS=Anopheles gambiae GN=AGAP006599 PE=4 SV=4
CCG018197.1	1861	466	2355	10.7597508	44.23790532	2.039638404	Up	0	0	AAEL002658-PA OS=Aedes aegypti GN=AAEL002658 PE=4 SV=1
CCG006124.1	28961	701	2356	1.04008094	2.843883156	1.451166394	Up	1.08E-12	2.64E-12	AAEL013441-PA OS=Aedes aegypti GN=TOLL9A PE=4 SV=2
CCG011173.1	1323	338	2357	10.9779065	62.28016504	2.504169811	Up	1.44E-13	4.32E-13	AAEL008375-PA OS=Aedes aegypti GN=AAEL008375 PE=4 SV=1
CCG009657.1	1957	586	2357	12.8667677	42.10355562	1.710292393	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_18280 PE=4 SV=1
CCG018322.1	1653	247	2359	6.42076494	49.88903503	2.957905675	Up	0	0	AAEL003676-PA OS=Aedes aegypti GN=AAEL003676 PE=4 SV=1
CCG022155.1	6846	665	2359	4.17395173	12.04595018	1.529062377	Up	3.93E-13	1.05E-12	AAEL013561-PA OS=Aedes aegypti GN=AAEL013561 PE=4 SV=1
CCG012744.1	3738	690	2359	7.93181297	22.06168403	1.475820355	Up	8.73E-13	2.18E-12	AAEL004984-PA OS=Aedes aegypti GN=AAEL004984 PE=4 SV=1
CCG008227.1	1077	34	2361	1.35651901	76.63555382	5.820032689	Up	8.82E-14	2.80E-13	AAEL003179-PA OS=Aedes aegypti GN=AAEL003179 PE=4 SV=1
CCG005379.2	3696	50	2361	0.58130052	22.3313018	5.26363934	Up	0	0	AAEL003893-PA OS=Aedes aegypti GN=AAEL003893 PE=4 SV=1

CCG022890.1	6393	555	2362	3.73036176	12.9159158	1.791762492	Up	0	0	AAEL001467-PA (Fragment) OS=Aedes aegypti GN=AAEL001467 PE=4 SV=1
CCG003215.3	3420	11027	2364	138.545983	24.16414219	-2.519425201	Down	0	0	6-phosphofructokinase OS=Aedes aegypti GN=AAEL006895 PE=3 SV=1
CCG015077.1	2043	179	2365	3.76484704	40.46809817	3.426121895	Up	0	0	in-phase inducer phosphatase 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001662 PE=4 SV=1
CCG006226.1	1875	528	2365	12.1002773	44.09403977	1.865543553	Up	0	0	AAEL011778-PA OS=Aedes aegypti GN=AAEL011778 PE=4 SV=1
CCG011453.1	9861	775	2366	3.37709607	8.387717558	1.312495062	Up	0	0	AAEL012471-PA OS=Aedes aegypti GN=DOME PE=4 SV=1
CCG022905.1	690	47533	2367	2960.11651	119.9220886	-4.625484644	Down	0	0	AAEL011184-PA OS=Aedes aegypti GN=AAEL011184 PE=3 SV=1
CCG008928.1	1707	554	2370	13.9456549	48.53609605	1.799242382	Up	9.41E-14	2.97E-13	AAEL001015-PA OS=Aedes aegypti GN=AAEL001015 PE=4 SV=1
CCG021750.1	6147	726	2370	5.07500038	13.47830095	1.40915881	Up	2.70E-13	7.54E-13	AAEL002392-PA OS=Aedes aegypti GN=AAEL002392 PE=4 SV=1
CCG027362.1	3306	772	2370	10.0340699	25.06083362	1.32052751	Up	2.17E-12	5.16E-12	AAEL006917-PA OS=Aedes aegypti GN=AAEL006917 PE=4 SV=1
CCG025493.1	3084	433	2372	6.03303991	26.88749433	2.155978284	Up	9.43E-13	2.34E-12	AAEL005455-PA OS=Aedes aegypti GN=AAEL005455 PE=4 SV=1
CCG019382.1	3660	506	2372	5.94062451	22.65601981	1.931207924	Up	3.38E-13	9.20E-13	AAEL013888-PA OS=Aedes aegypti GN=AAEL013888 PE=4 SV=1
CCG003747.1	2061	493	2373	10.278544	40.25035943	1.969365753	Up	6.97E-14	2.27E-13	AAEL011637-PA OS=Aedes aegypti GN=AAEL011637 PE=4 SV=1
CCG024025.1	2715	352	2374	5.57103005	30.56756871	2.455985805	Up	5.16E-13	1.34E-12	AAEL007558-PA (Fragment) OS=Aedes aegypti GN=AAEL007558 PE=4 SV=1
CCG000575.1	3122	531	2376	7.3084334	26.60501781	1.864064274	Up	2.40E-13	6.77E-13	AAEL003968-PA OS=Aedes aegypti GN=AAEL003968 PE=4 SV=1
CCG015416.1	1245	309	2377	10.6647775	66.74363364	2.645776364	Up	6.25E-13	1.59E-12	AAEL013085-PA OS=Aedes aegypti GN=AAEL013085 PE=4 SV=1

CCG020797.1	2617	595	2378	9.76958047	31.76567908	1.701100345	Up	7.20E-13	1.81E-12	AAEL009487-PA OS=Aedes aegypti GN=AAEL009487 PE=4 SV=1
CCG006357.1	2587	170	2380	2.82367796	32.16107411	3.509668126	Up	3.53E-13	9.52E-13	AAEL008706-PA OS=Aedes aegypti GN=AAEL008706 PE=4 SV=1
CCG015129.1	4835	922	2380	8.19402178	17.20800387	1.070436122	Up	0	0	AAEL001516-PA OS=Aedes aegypti GN=AAEL001516 PE=4 SV=1
CCG010135.1	2142	170	2383	3.4102964	38.89149092	3.511485504	Up	3.53E-13	9.52E-13	OrfGdpeptide transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ002172 PE=3 SV=1
CCG003853.1	47354	4710	2384	4.27392512	1.759947034	-1.28002962	Down	1.16E-291	7.26E-291	AAEL006973-PA OS=Aedes aegypti GN=AAEL006973 PE=4 SV=1
CCG000584.1	3897	358	2388	3.94743777	21.42170001	2.440084548	Up	1.30E-13	3.97E-13	AAEL006512-PA OS=Aedes aegypti GN=AAEL006512 PE=4 SV=1
CCG020616.1	1508	30941	2391	881.648912	55.42787783	-3.991520586	Down	0	0	Phosphoglycerate kinase OS=Aedes aegypti GN=PGK PE=3 SV=1
CCG005914.1	1479	50	2395	1.45266175	56.60924467	5.284266955	Up	0	0	AAEL012708-PA OS=Aedes aegypti GN=AAEL012708 PE=4 SV=1
CCG028227.1	1074	80	2397	3.20072511	78.02140542	4.607399303	Up	0	0	AAEL000387-PA OS=Aedes aegypti GN=AAEL000387 PE=4 SV=1
CCG001434.1	13314	765	2399	2.46896853	6.2990015	1.351214709	Up	0	0	AAEL010074-PA OS=Aedes aegypti GN=AAEL010074 PE=4 SV=1
CCG000825.1	1821	54	2400	1.27422607	46.0735114	5.176244392	Up	0	0	AAEL007176-PA OS=Aedes aegypti GN=AAEL007176 PE=3 SV=1
CCG002452.1	4406	511	2401	4.98355297	19.0501186	1.934553411	Up	0	0	AAEL005557-PA OS=Aedes aegypti GN=AAEL005557 PE=4 SV=1
CCG018908.1	1191	622	2402	22.4409529	70.5035943	1.651562869	Up	0	0	AAEL007422-PA OS=Aedes aegypti GN=AAEL007422 PE=4 SV=1
CCG017377.1	4254	194	2404	1.95959768	19.75545307	3.333621542	Up	5.97E-13	1.53E-12	AAEL010414-PA OS=Aedes aegypti GN=AAEL010414 PE=4 SV=1
CCG018013.1	827	506	2412	26.2910347	101.958118	1.955333821	Up	3.38E-13	9.20E-13	Nucleolar protein NOP5 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014566 PE=4 SV=1

CCG026040.3	2442	693	2415	12.1941139	34.5717602	1.503409135	Up	1.99E-12	4.75E-12	77 kDa echinoderm microtubule-associated protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005752 PE=4 SV=1
CCG006266.1	1754	5163	2415	126.483888	48.13240502	-1.393873205	Down	0	0	-
CCG023384.1	1857	183	2418	4.23449727	45.51917783	3.426211895	Up	3.05E-13	8.41E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_16294 PE=4 SV=1
CCG012913.1	1288	315	2419	10.5089025	65.65533502	2.643300239	Up	4.28E-13	1.13E-12	AAEL013914-PA OS=Aedes aegypti GN=AAEL013914 PE=4 SV=1
CCG028332.1	2183	215	2421	4.23201692	38.76957768	3.195507718	Up	0	0	AAEL000004-PA OS=Aedes aegypti GN=AAEL000004 PE=4 SV=1
CCG024806.1	603	743	2423	52.9461241	140.4708203	1.407673493	Up	0	0	AAEL012996-PA OS=Aedes aegypti GN=AAEL012996 PE=2 SV=1
CCG000097.1	3339	913	2424	11.7494363	25.37851539	1.111016137	Up	0	0	AGAP004049-PA OS=Anopheles gambiae GN=AgaP_AGAP004049 PE=4 SV=5
CCG016546.1	5936	4113	2426	29.7733353	14.2871933	-1.059298318	Down	1.53E-188	9.06E-188	AAEL005886-PA OS=Aedes aegypti GN=AAEL005886 PE=4 SV=1
CCG021294.1	3075	181	2431	2.52927544	27.6369336	3.449801495	Up	1.53E-13	4.55E-13	AAEL013487-PA OS=Aedes aegypti GN=AAEL013487 PE=4 SV=1
CCG005154.1	1512	357	2432	10.1456318	56.22918592	2.470460453	Up	3.04E-13	8.39E-13	Nuclear protein nops6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003353 PE=4 SV=1
CCG018294.1	1125	549	2432	20.9692305	75.57202588	1.849578378	Up	2.32E-13	6.62E-13	AAEL012925-PA OS=Aedes aegypti GN=AAEL012925 PE=4 SV=1
CCG026770.4	2832	200	2433	3.03458578	30.03301108	3.306977619	Up	0	0	DNA-binding protein smuap-z OS=Culex quinquefasciatus GN=CpipJ_CPIJ012615 PE=4 SV=1
CCG024065.1	1587	293	2435	7.93329064	53.63793569	2.757262406	Up	4.19E-13	1.11E-12	AAEL014904-PA OS=Aedes aegypti GN=AAEL014904 PE=3 SV=1
CCG006459.1	569	11993	2438	905.687218	149.7860787	-2.596109353	Down	0	0	NADH dehydrogenase [ubiquinone] 1 subunit C2 OS=Aedes aegypti GN=AAEL005946 PE=2 SV=1

CCG023777.1	3003	672	2440	9.61560495	28.40432745	1.562661213	Up	0	0	AAEL001636-PA OS=Aedes aegypti GN=AAEL001636 PE=4 SV=1
CCG016862.1	1647	208	2442	5.42665744	51.83249051	3.255720971	Up	3.93E-13	1.05E-12	AAEL003670-PA OS=Aedes aegypti GN=AAEL003670 PE=4 SV=1
CCG015361.1	3637	326	2443	3.85156268	23.48173499	2.608023198	Up	6.17E-13	1.58E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009453 PE=4 SV=1
CCG017137.1	2058	976	2443	20.3782609	41.49809045	1.026014015	Up	2.52E-12	5.94E-12	Probable transport protein Sec61 alpha subunit OS=Aedes aegypti GN=Sec61 PE=2 SV=1
CCG007657.1	3030	863	2446	12.2385746	28.22044389	1.205305143	Up	0	0	AAEL006032-PA OS=Aedes aegypti GN=AAEL006032 PE=4 SV=1
CCG019981.1	4875	4900	2451	43.1900922	17.57594592	-1.297098061	Down	0	0	AAEL005590-PA OS=Aedes aegypti GN=AAEL005590 PE=4 SV=1
CCG018804.1	840	656	2452	33.5573166	102.0448746	1.604504463	Up	9.98E-13	2.47E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_00971 PE=4 SV=1
CCG015793.1	1116	759	2452	29.2240399	76.80797011	1.394100392	Up	1.54E-12	3.70E-12	AAEL005655-PA OS=Aedes aegypti GN=AAEL005655 PE=4 SV=1
CCG020572.1	3084	700	2453	9.7531823	27.80565918	1.511433611	Up	0	0	Subunitase OS=Culex quinquefasciatus GN=CpipJ_CPIJ002113 PE=4 SV=1
CCG005662.1	768	19	2454	1.06305333	111.7026187	6.715305224	Up	7.73E-10	1.60E-09	AAEL013338-PA OS=Aedes aegypti GN=AAEL013338 PE=3 SV=1
CCG003977.1	3246	130	2454	1.72090742	26.42871571	3.940864924	Up	0	0	AAEL009738-PA OS=Aedes aegypti GN=AAEL009738 PE=4 SV=1
CCG022210.1	1722	390	2455	9.73182143	49.83889052	2.356490199	Up	6.59E-13	1.68E-12	AAEL000905-PA (Fragment) OS=Aedes aegypti GN=AAEL000905 PE=4 SV=2
CCG028346.1	2115	940	2455	19.0976598	40.57804703	1.087303566	Up	0	0	AAEL012746-PA OS=Aedes aegypti GN=AAEL012746 PE=3 SV=1
CCG012942.1	808	4284	2458	227.824682	106.3458469	-1.099160361	Down	2.05E-208	1.23E-207	AAEL001969-PA OS=Aedes aegypti GN=AAEL001969 PE=4 SV=1

CCG011685.1	2267	72	2459	1.36472029	37.91901305	4.796244222	Up	0	0	DNA repair and recombination protein pif1, mitochondrial OS=Culex quinquefasciatus GN=CpipJ_CPIJ004811 PE=4 SV=1
CCG011942.1	2976	285	2459	4.11504515	28.88521592	2.811351114	Up	4.57E-13	1.20E-12	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015487 PE=4 SV=1
CCG023976.1	5973	380	2461	2.73371826	14.40353577	2.397486538	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006203 PE=4 SV=1
CCG022359.1	1233	222	2462	7.73664321	69.8031447	3.173512384	Up	0	0	AAEL001495-PA OS=Aedes aegypti GN=AAEL001495 PE=4 SV=1
CCG017127.1	7374	494	2464	2.87863424	11.68120341	2.020732513	Up	0	0	AAEL008398-PA OS=Aedes aegypti GN=AAEL008398 PE=4 SV=1
CCG018628.1	39122	5962	2465	6.54837579	2.202652018	-1.571895525	Down	0	0	AAEL012064-PA OS=Aedes aegypti GN=AAEL012064 PE=4 SV=1
CCG004309.1	1733	4583	2466	113.635484	49.74443769	-1.191806282	Down	1.13E-253	6.99E-253	AAEL010148-PA OS=Aedes aegypti GN=AAEL010148 PE=3 SV=1
CCG006251.1	3723	4626	2468	53.3918862	23.17406046	-1.204109668	Down	3.09E-260	1.91E-259	AAEL010585-PA OS=Aedes aegypti GN=AAEL010585 PE=3 SV=1
CCG018042.1	9653	283	2469	1.25975706	8.941467456	2.827366082	Up	0	0	AAEL012187-PA OS=Aedes aegypti GN=AAEL012187 PE=4 SV=1
CCG005204.1	3591	492	2470	5.88724852	24.04537556	2.030094025	Up	0	0	AAEL006415-PA OS=Aedes aegypti GN=AAEL006415 PE=3 SV=1
CCG014425.2	13592	12044	2470	38.075889	6.3527769	-2.583418462	Down	0	0	AAEL009232-PA OS=Aedes aegypti GN=AAEL009232 PE=4 SV=1
CCG022162.1	2012	751	2474	16.0389019	42.98547551	1.422273891	Up	0	0	AAEL013553-PA OS=Aedes aegypti GN=AAEL013553 PE=4 SV=1
CCG007620.3	2231	798	2474	15.3697213	38.76592413	1.334698053	Up	1.86E-12	4.46E-12	AAEL010372-PA OS=Aedes aegypti GN=AAEL010372 PE=4 SV=1
CCG001863.1	1684	189	2476	4.82261273	51.39946157	3.413866379	Up	2.34E-13	6.64E-13	AAEL008220-PA OS=Aedes aegypti GN=AAEL008220 PE=4 SV=1
CCG024530.1	7848	346	2476	1.89443529	11.02914033	2.541480575	Up	2.24E-13	6.40E-13	AAEL005814-PA OS=Aedes aegypti GN=AAEL005814 PE=4 SV=1

CCG000894.1	3624	270	2478	3.20138751	23.90358991	2.900458079	Up	2.59E-13	7.26E-13	AAEL001436-PA OS=Aedes aegypti GN=AAEL001436 PE=4 SV=1
CCG003867.1	11292	16052	2478	61.0830836	7.67150282	-2.993191765	Down	0	0	AAEL002436-PA (Fragment) OS=Aedes aegypti GN=AAEL002436 PE=2 SV=1
CCG023083.1	32125	900	2479	1.20382136	2.697636362	1.164074569	Up	4.28E-13	1.13E-12	AAEL014816-PA OS=Aedes aegypti GN=AAEL014816 PE=4 SV=1
CCG018345.1	3018	463	2480	6.59210972	28.72648323	2.123569226	Up	6.16E-13	1.57E-12	AGAP000894-PA OS=Anopheles gambiae GN=AgaP_AGAP000894 PE=4 SV=5
CCG028425.2	3559	489	2482	5.90396185	24.37944449	2.045909949	Up	1.91E-13	5.52E-13	AAEL001416-PA OS=Aedes aegypti GN=AAEL001416 PE=4 SV=1
CCG006512.1	816	144	2483	7.58289434	106.3742662	3.810256749	Up	7.53E-14	2.43E-13	AAEL009235-PA OS=Aedes aegypti GN=AAEL009235 PE=4 SV=1
CCG015459.1	2612	312	2483	5.13267887	33.23177689	2.694779531	Up	9.28E-14	2.94E-13	AAEL003388-PA OS=Aedes aegypti GN=AAEL003388 PE=4 SV=1
CCG015567.1	29822	545	2486	0.78527615	2.91416659	1.891811365	Up	0	0	AAEL008468-PA OS=Aedes aegypti GN=AAEL008468 PE=4 SV=1
CCG014115.1	1475	850	2489	24.7622199	58.99061077	1.252344688	Up	2.37E-12	5.61E-12	AAEL001255-PA OS=Aedes aegypti GN=AAEL001255 PE=4 SV=1
CCG020010.1	3069	368	2491	5.15244781	28.37441103	2.461260554	Up	3.69E-13	9.91E-13	AAEL003108-PA OS=Aedes aegypti GN=AAEL003108 PE=4 SV=1
CCG020055.1	2211	649	2492	12.6130067	39.40118757	1.643326889	Up	1.65E-12	3.96E-12	AGAP002832-PA OS=Anopheles gambiae GN=AgaP_AGAP002832 PE=4 SV=5
CCG010237.1	898	179	2495	8.56523663	97.12795161	3.503321527	Up	0	0	AAEL007951-PA OS=Aedes aegypti GN=GSTe2 PE=2 SV=1
CCG014843.2	4796	401	2495	3.59275721	18.18617609	2.339678878	Up	1.05E-13	3.28E-13	AAEL010022-PA OS=Aedes aegypti GN=AAEL010022 PE=4 SV=1
CCG014938.1	3024	639	2495	9.07991416	28.84289039	1.667465183	Up	4.28E-13	1.13E-12	AAEL011258-PA OS=Aedes aegypti GN=AAEL011258 PE=4 SV=1
CCG003591.1	4611	164	2496	1.5283098	18.92341332	3.630163418	Up	1.14E-13	3.54E-13	AAEL006211-PA (Fragment) OS=Aedes aegypti GN=AAEL006211 PE=4 SV=1

CCG020219.1	3198	403	2496	5.41488526	27.2845087	2.333079394	Up	0	0	AAEL001372-PA OS=Aedes aegypti GN=AAEL001372 PE=4 SV=1
CCG023315.1	1771	429	2496	10.4088177	49.26925964	2.242881585	Up	0	0	AAEL002867-PA OS=Aedes aegypti GN=AAEL002867 PE=3 SV=1
CCG019489.1	21627	5390	2498	10.7091538	4.037812705	-1.407198592	Down	0	0	AAEL001056-PA OS=Aedes aegypti GN=AAEL001056 PE=4 SV=1
CCG017528.1	1069	836	2500	33.6040207	81.75462295	1.282666451	Up	1.21E-12	2.95E-12	AAEL010032-PA (Fragment) OS=Aedes aegypti GN=AAEL010032 PE=2 SV=1
CCG016892.1	885	11534	2502	560.014598	98.83119603	-2.502426029	Down	0	0	AAEL009906-PA OS=Aedes aegypti GN=AAEL009906 PE=4 SV=1
CCG014419.1	3572	692	2503	8.32448386	24.49623929	1.557127552	Up	0	0	AAEL004727-PA OS=Aedes aegypti GN=AAEL004727 PE=4 SV=1
CCG019775.1	13644	744	2503	2.34311658	6.413116884	1.452596968	Up	0	0	AAEL007041-PA OS=Aedes aegypti GN=AAEL007041 PE=4 SV=1
CCG012797.1	1317	8516	2503	277.851374	66.43930658	-2.064204455	Down	0	0	AAEL010330-PA OS=Aedes aegypti GN=AAEL010330 PE=4 SV=1
CCG028381.1	2615	75	2506	1.23240157	33.50112489	4.764665213	Up	0	0	AAEL013544-PA OS=Aedes aegypti GN=AAEL013544 PE=4 SV=1
CCG022984.1	4066	350	2506	3.69882123	21.54585381	2.542272791	Up	3.00E-14	1.04E-13	AAEL014265-PA OS=Aedes aegypti GN=AAEL014265 PE=4 SV=1
CCG007158.1	4734	813	2507	7.37946646	18.51296998	1.326947943	Up	0	0	Cuniflexin43-interacting protein of 150 kDa OS=Culex quinquefasciatus GN=CpipJ_CPIJ015283 PE=4 SV=1
CCG006587.2	1158	854	2507	31.6892516	75.68255602	1.255967225	Up	1.02E-12	2.51E-12	AAEL000926-PA OS=Aedes aegypti GN=AAEL008928 PE=3 SV=1
CCG000133.1	2718	804	2511	12.7106941	32.29589146	1.345307826	Up	2.72E-13	7.60E-13	Fluoperin OS=Culex quinquefasciatus GN=CpipJ_CPIJ004052 PE=4 SV=1
CCG000618.1	714	371	2514	22.3274111	123.0883863	2.462806762	Up	5.41E-13	1.39E-12	AAEL000675-PA OS=Aedes aegypti GN=AAEL006751 PE=4 SV=1
CCG008925.1	2130	608	2514	12.2655393	41.26061399	1.750154625	Up	1.30E-12	3.16E-12	AAEL000572-PA OS=Aedes aegypti GN=AAEL000572 PE=4 SV=1

CCG019373.1	1683	418	2518	10.6722217	52.30240102	2.293016639	Up	6.74E-13	1.70E-12	AAEL007882-PA OS=Aedes aegypti GN=AAEL007882 PE=4 SV=1
CCG011410.1	15163	682	2519	1.93268872	5.807551223	1.587320682	Up	1.08E-12	2.66E-12	AAEL009637-PA OS=Aedes aegypti GN=CAT-B3 PE=2 SV=1
CCG027641.1	2217	97	2520	1.88004703	39.73606561	4.40160838	Up	3.62E-14	1.25E-13	AAEL012725-PA OS=Aedes aegypti GN=AAEL012725 PE=4 SV=1
CCG012951.1	50139	4668	2521	4.00053294	1.757709881	-1.186495239	Down	1.65E-256	1.02E-255	AAEL006573-PA OS=Aedes aegypti GN=AAEL006573 PE=4 SV=1
CCG002115.1	3342	172	2522	2.21148844	26.38084202	3.576401009	Up	1.35E-13	4.09E-13	AAEL010857-PA OS=Aedes aegypti GN=AAEL010857 PE=4 SV=1
CCG014226.1	10046	524	2524	2.24130409	8.783066949	1.970386397	Up	0	0	AAEL003887-PA OS=Aedes aegypti GN=AAEL003887 PE=4 SV=1
CCG008769.1	1563	131	2528	3.60143009	56.54160184	3.97267095	Up	1.26E-13	3.87E-13	AAEL010437-PA (Fragment) OS=Aedes aegypti GN=AAEL010437 PE=4 SV=1
CCG001795.1	1056	595	2530	24.2111668	83.75420477	1.790489015	Up	7.20E-13	1.81E-12	AAEL002487-PA OS=Aedes aegypti GN=AAEL002487 PE=2 SV=1
CCG013400.1	1663	87	2531	2.24796567	53.20468942	4.5648615	Up	0	0	AAEL009654-PA OS=Aedes aegypti GN=AAEL009654 PE=4 SV=1
CCG015929.1	2514	298	2531	5.09346894	35.19466926	2.788636475	Up	3.92E-13	1.05E-12	Ribonucleoside-diphosphate reductase OS=Aedes aegypti GN=RNR1 PE=2 SV=1
CCG020998.1	1161	468	2532	17.3211333	76.23975606	2.138010174	Up	5.00E-13	1.30E-12	AAEL012955-PA OS=Aedes aegypti GN=AAEL012955 PE=4 SV=1
CCG015799.1	5772	388	2535	2.88847142	15.35329723	2.410170394	Up	3.43E-13	9.27E-13	AAEL001361-PA (Fragment) OS=Aedes aegypti GN=AAEL001361 PE=4 SV=1
CCG026595.1	3228	821	2535	10.9287956	27.45329356	1.328844824	Up	0	0	AAEL007329-PA OS=Aedes aegypti GN=AAEL007329 PE=4 SV=1
CCG028390.1	19467	48	2536	0.10595096	4.55407561	5.425689733	Up	5.35E-14	1.79E-13	AAEL009993-PA OS=Aedes aegypti GN=AAEL009993 PE=4 SV=1
CCG016078.1	1527	415	2538	11.6780877	58.10354057	2.314822031	Up	0	0	AAEL000081-PA (Fragment) OS=Aedes aegypti GN=AAEL000081 PE=3 SV=1

CCG019024.1	2555	359	2539	6.03762612	34.73935997	2.52451785	Up	0	0	mystery 45A OS=Culex quinquefasciatus GN=CpipJ_CPIJ001550 PE=4 SV=1
CCG003809.1	1830	5056	2541	118.718567	48.54042693	-1.290286884	Down	0	0	AAEL013253-PA OS=Aedes aegypti GN=AAEL013253 PE=4 SV=2
CCG015650.1	1587	284	2545	7.68960594	56.06100465	2.866016025	Up	0	0	AAEL003371-PA OS=Aedes aegypti GN=AAEL003371 PE=4 SV=1
CCG010633.1	1512	420	2545	11.9360374	58.84180846	2.301517627	Up	1.84E-13	5.32E-13	Nuclear protein nops6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003353 PE=4 SV=1
CCG025662.1	2002	485	2545	10.4097509	44.43996723	2.093922208	Up	2.07E-13	5.94E-13	AAEL001339-PA OS=Aedes aegypti GN=AAEL001339 PE=4 SV=1
CCG018470.1	6792	374	2547	2.36611907	13.10935379	2.470001988	Up	1.37E-13	4.12E-13	AAEL010978-PA OS=Aedes aegypti GN=AAEL010978 PE=4 SV=1
CCG012875.1	3714	930	2549	10.7597881	23.9926353	1.156941956	Up	0	0	Neuropathy target esterase/swiss cheese OS=Culex quinquefasciatus GN=CpipJ_CPIJ003198 PE=4 SV=1
CCG023120.1	1848	8297	2550	192.922017	48.23788191	-1.999779335	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008688 PE=4 SV=1
CCG009839.1	6030	172	2552	1.22567071	14.79494566	3.593461063	Up	1.35E-13	4.09E-13	AAEL010499-PA (Fragment) OS=Aedes aegypti GN=AAEL010499 PE=4 SV=1
CCG015997.1	1738	557	2554	13.771083	51.37136874	1.899322496	Up	5.29E-13	1.37E-12	AAEL012973-PA OS=Aedes aegypti GN=AAEL012973 PE=4 SV=1
CCG026396.1	2108	891	2554	18.162255	42.35457252	1.221574392	Up	0	0	AAEL002374-PA OS=Aedes aegypti GN=AAEL002374 PE=4 SV=1
CCG028025.1	2619	32	2558	0.52502158	34.14405192	6.023113753	Up	1.77E-13	5.17E-13	AAEL013025-PA OS=Aedes aegypti GN=AAEL013025 PE=4 SV=1
CCG015415.1	3523	348	2558	4.24452672	25.38270564	2.580170257	Up	0	0	AAEL013086-PA OS=Aedes aegypti GN=AAEL013086 PE=4 SV=1
CCG021780.1	3246	758	2559	10.0342141	27.55952873	1.457623598	Up	0	0	AAEL009061-PA OS=Aedes aegypti GN=AAEL009061 PE=4 SV=1

CCG024298.1	6692	577	2560	3.70495171	13.37316027	1.85181379	Up	4.04E-13	1.07E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013516 PE=4 SV=1
CCG002029.1	2599	863	2560	14.2681343	34.43370086	1.27102455	Up	0	0	AAEL009883-PA OS=Aedes aegypti GN=AAEL009883 PE=4 SV=1
CCG025342.1	971	14423	2560	638.262083	92.16600261	-2.791842359	Down	0	0	AAEL008002-PA OS=Aedes aegypti GN=AAEL008002 PE=4 SV=1
CCG004233.1	11053	202	2561	0.78529688	8.099895668	3.366593259	Up	0	0	AAEL008123-PA OS=Aedes aegypti GN=AAEL008123 PE=4 SV=1
CCG021318.1	752	977	2561	55.8263706	119.0533867	1.092589989	Up	2.38E-12	5.63E-12	AAEL008039-PA OS=Aedes aegypti GN=AAEL008039 PE=4 SV=1
CCG010096.1	4669	725	2562	6.67231904	19.1825027	1.523530779	Up	0	0	AAEL011145-PA (Fragment) OS=Aedes aegypti GN=AAEL011145 PE=4 SV=1
CCG023879.1	2237	208	2565	3.99539776	40.08403215	3.326616596	Up	3.93E-13	1.05E-12	AAEL000001-PA OS=Aedes aegypti GN=AAEL000001 PE=4 SV=1
CCG004901.1	628	11800	2566	807.393103	142.8390736	-2.49888058	Down	0	0	AAEL000986-PB OS=Aedes aegypti GN=AAEL000986 PE=4 SV=1
CCG003382.1	2741	166	2567	2.60232614	32.73910853	3.653141353	Up	1.83E-13	5.32E-13	AAEL014051-PA OS=Aedes aegypti GN=AAEL014051 PE=4 SV=1
CCG013153.1	1164	141	2569	5.20509672	77.15447855	3.889753026	Up	0	0	AAEL004455-PA OS=Aedes aegypti GN=AAEL004455 PE=4 SV=1
CCG010228.1	1440	218	2570	6.50514038	62.39081341	3.261681523	Up	0	0	AAEL006205-PA OS=Aedes aegypti GN=AAEL006205 PE=4 SV=1
CCG028444.1	3035	856	2572	12.1193057	29.62526783	1.289521145	Up	1.31E-12	3.17E-12	AAEL014540-PA OS=Aedes aegypti GN=AAEL014540 PE=4 SV=1
CCG014556.2	2498	288	2573	4.95407669	36.00786475	2.861623944	Up	0	0	AAEL001587-PA OS=Aedes aegypti GN=AAEL001587 PE=4 SV=1
CCG021880.1	1293	215	2574	7.14500614	69.59211478	3.283916692	Up	0	0	AAEL012235-PA OS=Aedes aegypti GN=AAEL012235 PE=2 SV=1
CCG023187.1	869	257	2580	12.7079652	103.7886698	3.029844005	Up	4.14E-13	1.10E-12	AAEL009667-PA OS=Aedes aegypti GN=AAEL009667 PE=4 SV=1
CCG015663.1	3285	535	2580	6.99811507	27.45581555	1.972073473	Up	2.42E-14	8.44E-14	AAEL003901-PA OS=Aedes aegypti GN=AAEL003901 PE=4 SV=1

CCG008411.1	2438	898	2584	15.8272443	37.05175848	1.227131924	Up	2.22E-12	5.28E-12	AAEL003083-PA OS=Aedes aegypti GN=AAEL003083 PE=4 SV=1
CCG012878.1	3747	183	2585	2.09860193	24.11719921	3.522561931	Up	3.05E-13	8.41E-13	AAEL001238-PA OS=Aedes aegypti GN=AAEL001238 PE=4 SV=1
CCG021018.3	5940	357	2586	2.58252445	15.21920938	2.5590395	Up	3.04E-13	8.39E-13	AAEL010633-PA OS=Aedes aegypti GN=AAEL010633 PE=4 SV=1
CCG010205.1	4419	447	2586	4.34656514	20.45759306	2.234688742	Up	3.44E-13	9.29E-13	THO complex subunit 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ019403 PE=4 SV=1
CCG026416.1	2716	600	2586	9.4925776	33.2850161	1.810001073	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_01164 PE=4 SV=1
CCG001311.1	4620	235	2588	2.18568996	19.58268837	3.163418159	Up	4.34E-13	1.14E-12	AAEL000924-PA OS=Aedes aegypti GN=AAEL000924 PE=4 SV=1
CCG023468.1	3582	367	2589	4.40253841	25.26716319	2.520856201	Up	4.12E-13	1.09E-12	DNA polymerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ018287 PE=3 SV=1
CCG012990.1	3567	422	2592	5.08360751	25.40281845	2.321064018	Up	5.52E-13	1.42E-12	AAEL001651-PA OS=Aedes aegypti GN=AAEL001651 PE=4 SV=1
CCG011145.1	957	605	2593	27.1647748	94.71976141	1.801928363	Up	2.07E-13	5.93E-13	AAEL000950-PA OS=Aedes aegypti GN=AAEL000950 PE=4 SV=1
CCG028138.1	2312	215	2594	3.99588795	39.22221884	3.295083118	Up	0	0	AAEL001681-PA OS=Aedes aegypti GN=AAEL001681 PE=4 SV=1
CCG028452.1	3329	610	2595	7.87369724	27.25044404	1.791166595	Up	3.62E-13	9.71E-13	AAEL012019-PA OS=Aedes aegypti GN=AAEL012019 PE=4 SV=1
CCG004043.2	995	714	2596	30.8345633	91.20772513	1.564607608	Up	1.92E-13	5.54E-13	AAEL000932-PA OS=Aedes aegypti GN=AAEL000932 PE=4 SV=1
CCG016557.1	2382	129	2597	2.32707631	38.11362081	4.033716247	Up	0	0	AAEL003071-PA OS=Aedes aegypti GN=AAEL003071 PE=4 SV=1
CCG013765.1	1939	346	2603	7.6676267	46.92954845	2.613644573	Up	2.24E-13	6.39E-13	AAEL011341-PA OS=Aedes aegypti GN=AAEL011341 PE=3 SV=1
CCG027288.1	2140	584	2603	11.7263201	42.52167964	1.858448241	Up	3.10E-13	8.52E-13	AAEL000996-PA OS=Aedes aegypti GN=AAEL000996 PE=4 SV=1
CCG024931.1	5133	968	2603	8.10339044	17.7277215	1.129409563	Up	7.03E-13	1.77E-12	AAEL000152-PA OS=Aedes aegypti GN=AAEL006152 PE=4 SV=1

CCG006992.1	1869	115	2604	2.64393766	48.70591371	4.203336886	Up	6.13E-14	2.03E-13	AAEL009508-PA OS=Aedes aegypti GN=AAEL009508 PE=4 SV=1
CCG015312.2	3517	295	2605	3.60422852	25.89317913	2.844809717	Up	2.32E-13	6.62E-13	AAEL001324-PA OS=Aedes aegypti GN=AAEL001324 PE=4 SV=1
CCG011899.1	1956	490	2605	10.7644018	46.55741871	2.112742922	Up	0	0	AAEL007329-PA OS=Aedes aegypti GN=AAEL007329 PE=4 SV=1
CCG008335.1	2222	1000	2607	19.3383144	41.01540394	1.084703787	Up	1.46E-12	3.53E-12	-
CCG004360.1	5415	592	2609	4.6977069	16.84323991	1.842141067	Up	4.13E-13	1.10E-12	AAEL008796-PA OS=Aedes aegypti GN=AAEL008796 PE=4 SV=1
CCG013455.2	4777	593	2609	5.33411192	19.09276619	1.839706138	Up	1.13E-12	2.78E-12	AAEL008482-PA OS=Aedes aegypti GN=AAEL008482 PE=4 SV=1
CCG020701.1	3195	253	2610	3.40261122	28.55746553	3.069153721	Up	3.13E-13	8.62E-13	AAEL003495-PA (Fragment) OS=Aedes aegypti GN=AAEL003495 PE=4 SV=1
CCG012628.1	687	479	2610	29.9599751	132.8109205	2.14826545	Up	0	0	AAEL012807-PA OS=Aedes aegypti GN=AAEL012807 PE=4 SV=1
CCG019913.1	2358	760	2610	13.849448	38.69427582	1.482291687	Up	1.99E-12	4.75E-12	AAEL010912-PA OS=Aedes aegypti GN=AAEL010912 PE=4 SV=1
CCG024143.1	5595	994	2615	7.63394391	16.33885501	1.097806393	Up	3.37E-12	7.85E-12	AAEL001141-PA OS=Aedes aegypti GN=AAEL001141 PE=4 SV=1
CCG025573.1	3762	664	2616	7.58423811	24.30910474	1.680420598	Up	4.96E-13	1.29E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003705 PE=4 SV=1
CCG027591.1	3441	8564	2616	106.943565	26.57682419	-2.008609051	Down	0	0	AAEL012556-PA OS=Aedes aegypti GN=AAEL012556 PE=4 SV=1
CCG016252.1	5010	513	2617	4.39989498	18.26064078	2.053196397	Up	0	0	Rho GTPase-activating protein 5 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009693 PE=4 SV=1
CCG012754.1	5073	419	2618	3.54904766	18.04075864	2.345756152	Up	0	0	AAEL012935-PA OS=Aedes aegypti GN=AAEL012935 PE=4 SV=1
CCG007080.1	3705	673	2618	7.80529862	24.70196183	1.662099891	Up	7.17E-14	2.33E-13	Ubiquitin carboxyl-terminal hydrolase OS=Aedes aegypti GN=AAEL009702 PE=3 SV=1
CCG024306.1	2742	858	2620	13.4456719	33.40287569	1.312830463	Up	1.49E-14	5.27E-14	AAEL013581-PA OS=Aedes aegypti GN=AAEL013581 PE=4 SV=1

CCG015780.1	1695	9009	2620	228.386041	54.03580244	-2.07948696	Down	0	0	Mitochondrial NADH:ubiquinone oxidoreductase NDUFV1/51 kDa subunit OS=Aedes aegypti PE=2 SV=1
CCG007531.1	1347	2	2622	0.06380065	68.04795969	10.05876517	Up	9.36E-05	0.0001535	AAEL004172-PA OS=Aedes aegypti GN=AAEL004172 PE=3 SV=1
CCG017444.1	4932	377	2624	3.28458839	18.59905074	2.501444495	Up	4.66E-13	1.21E-12	AAEL014774-PA OS=Aedes aegypti GN=AAEL014774 PE=4 SV=1
CCG015638.1	1137	457	2624	17.2710367	80.67767656	2.223814853	Up	1.13E-12	2.77E-12	AAEL008739-PA OS=Aedes aegypti GN=AAEL008739 PE=4 SV=1
CCG012836.1	19192	783	2628	1.75308994	4.786908678	1.449194267	Up	0	0	AAEL003540-PA OS=Aedes aegypti GN=AAEL003540 PE=4 SV=1
CCG008560.1	444	5	2634	0.48389341	207.3876149	8.743424739	Up	1.59E-05	2.76E-05	Histone H2B OS=Aedes aegypti GN=AAEL015682 PE=3 SV=1
CCG008598.1	3952	124	2634	1.34824066	23.2996207	4.111156524	Up	1.61E-13	4.77E-13	AAEL004286-PA OS=Aedes aegypti GN=AAEL004286 PE=4 SV=1
CCG016408.1	1734	555	2634	13.7532888	53.10271108	1.949008873	Up	0	0	AAEL010345-PA OS=Aedes aegypti GN=AAEL010345 PE=4 SV=1
CCG010675.1	1128	24909	2634	948.876879	81.63129523	-3.53902665	Down	0	0	AAEL002417-PA OS=Aedes aegypti GN=AAEL002417 PE=4 SV=1
CCG025450.1	14940	347	2635	0.9980253	6.165666619	2.627108598	Up	0	0	AAEL014053-PA OS=Aedes aegypti GN=AAEL014053 PE=4 SV=1
CCG019185.1	1299	235	2638	7.77358555	70.99302088	3.191025107	Up	4.34E-13	1.14E-12	AAEL013546-PA OS=Aedes aegypti GN=AAEL013546 PE=3 SV=1
CCG024509.1	4221	943	2638	9.59972986	21.84788773	1.186428092	Up	1.51E-12	3.64E-12	AAEL012325-PA OS=Aedes aegypti GN=AAEL012325 PE=4 SV=1
CCG020346.1	1327	611	2639	19.784859	69.52139593	1.813060269	Up	2.21E-13	6.34E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009391 PE=4 SV=1
CCG006929.1	888	59	2642	2.85497111	104.0087469	5.187084906	Up	0	0	AAEL008085-PA OS=Aedes aegypti GN=AAEL008085 PE=2 SV=1
CCG001162.2	1254	497	2642	17.0302696	73.65212698	2.112625914	Up	7.04E-14	2.29E-13	-
CCG014211.1	2469	1067	2642	18.5697476	37.40776316	1.010383494	Up	2.93E-12	6.87E-12	AAEL003489-PA (Fragment) OS=Aedes aegypti GN=AAEL003489 PE=4 SV=1

CCG019523.1	1888	868	2643	19.7551534	48.93788427	1.308722681	Up	0	0	AAEL006633-PA OS=Aedes aegypti GN=IAP2 PE=4 SV=2
CCG017209.1	3286	323	2644	4.22374446	28.12832738	2.735429311	Up	1.55E-13	4.61E-13	AAEL001497-PA (Fragment) OS=Aedes aegypti GN=AAEL001497 PE=4 SV=1
CCG017692.1	603	899	2644	64.0626723	153.2830577	1.25864236	Up	0	0	AAEL010765-PA OS=Aedes aegypti GN=AAEL010765 PE=4 SV=1
CCG014937.1	1574	455	2646	12.4213655	58.7672175	2.242187815	Up	0	0	AAEL011251-PA OS=Aedes aegypti GN=AAEL011251 PE=4 SV=1
CCG024496.1	5574	355	2649	2.73668026	16.61364822	2.601870118	Up	2.28E-13	6.52E-13	AAEL014260-PA OS=Aedes aegypti GN=AAEL014260 PE=4 SV=1
CCG001921.1	1645	759	2650	19.8261572	56.31576501	1.506133773	Up	1.54E-12	3.70E-12	AAEL001588-PA OS=Aedes aegypti GN=AAEL001588 PE=4 SV=1
CCG011356.1	2496	772	2650	13.2903186	37.11515763	1.481632811	Up	2.17E-12	5.15E-12	AAEL008686-PA OS=Aedes aegypti GN=AAEL008686 PE=4 SV=1
CCG010797.1	3390	64	2652	0.81122803	27.34789086	5.075178264	Up	0	0	DNA-directed RNA polymerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ002457 PE=3 SV=1
CCG018960.1	3315	490	2654	6.35148415	27.98771238	2.13962792	Up	0	0	AAEL012553-PA OS=Aedes aegypti GN=HOP PE=4 SV=2
CCG010326.1	23511	7553	2654	13.804194	3.946206735	-1.806568212	Down	0	0	AAEL014378-PA OS=Aedes aegypti GN=AAEL014378 PE=4 SV=1
CCG000614.1	2209	373	2655	7.25564102	42.01639875	2.533777529	Up	3.38E-13	9.19E-13	AAEL003377-PA OS=Aedes aegypti GN=AAEL003377 PE=4 SV=1
CCG024797.3	5194	480	2655	3.97101899	17.86950805	2.169918754	Up	5.41E-13	1.39E-12	AAEL015099-PA OS=Aedes aegypti GN=AAEL015099 PE=4 SV=1
CCG011352.1	905	2	2656	0.09496074	102.5957824	10.07735264	Up	9.36E-05	0.00015404	Cation efflux protein/ zinc transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ010101 PE=4 SV=1
CCG018953.1	1198	987	2656	35.4016094	77.50349174	1.130446361	Up	3.78E-12	8.77E-12	AAEL006480-PA OS=Aedes aegypti GN=AAEL006480 PE=4 SV=1
CCG025951.1	1758	62	2657	1.51542864	52.83512024	5.123699406	Up	0	0	AAEL009454-PA OS=Aedes aegypti GN=AAEL009454 PE=4 SV=1

CCG015969.1	1851	89	2658	2.06607584	50.19940554	4.602705162	Up	0	0	AAEL015216-PA OS=Aedes aegypti GN=AAEL015216 PE=3 SV=1
CCG019719.1	7034	648	2658	3.95854251	13.20999427	1.73858859	Up	0	0	AGAP007731-PA (Fragment) OS=Anopheles gambiae GN=AGAP007731 PE=4 SV=4
CCG023804.1	3264	222	2661	2.92257386	28.49999218	3.285650132	Up	0	0	AAEL011261-PA OS=Aedes aegypti GN=AAEL011261 PE=4 SV=1
CCG017517.1	1023	5414	2661	227.407765	90.93252638	-1.322413173	Down	0	0	Ca2+-binding protein Regucalcin/SMP30 OS=Aedes aegypti PE=2 SV=1
CCG014260.1	993	5707	2663	246.956974	93.75014204	-1.397366931	Down	0	0	AAEL010684-PA OS=Aedes aegypti GN=AAEL010684 PE=4 SV=1
CCG007772.1	3894	20508	2663	226.302855	23.90700849	-3.242749265	Down	0	0	AAEL009510-PA OS=Aedes aegypti GN=AAEL009510 PE=2 SV=1
CCG024200.1	606	534	2666	37.8644196	153.793343	2.022078337	Up	0	0	DNA-directed RNA polymerase subunit OS=Aedes aegypti GN=AAEL001831 PE=2 SV=1
CCG002169.1	681	851	2666	53.6963938	136.8557502	1.349758947	Up	1.91E-12	4.57E-12	AAEL005512-PA OS=Aedes aegypti GN=AAEL005512 PE=4 SV=1
CCG006767.1	564	390	2667	29.7131144	165.3080215	2.475985	Up	6.59E-13	1.68E-12	AAEL003552-PA OS=Aedes aegypti GN=AAEL003552 PE=4 SV=1
CCG028267.1	4890	240	2668	2.10894403	19.07335019	3.176965559	Up	0	0	Rothmund-Thomson syndrome DNA helicase recq4 OS=Culex quinquefasciatus GN=CpipJ_CPIJ007409 PE=4 SV=1
CCG001204.1	2571	317	2669	5.29809641	36.29079763	2.776057764	Up	2.22E-13	6.35E-13	AGAP004341-PA OS=Anopheles gambiae GN=AgaP_AGAP004341 PE=4 SV=5
CCG009683.1	1500	439	2669	12.575809	62.20242714	2.306319664	Up	0	0	AAEL001562-PA OS=Aedes aegypti GN=AAEL001562 PE=4 SV=1
CCG011759.1	2973	715	2670	10.3341272	31.39542515	1.603137799	Up	0	0	AAEL003375-PA OS=Aedes aegypti GN=AAEL003375 PE=4 SV=1
CCG003954.1	6537	6715	2670	44.139784	14.2785068	-1.628234454	Down	0	0	Gag polyprotein OS=Anopheles gambiae PE=4 SV=1

CCG028215.1	23899	6537	2671	11.7533435	3.907006873	-1.588935517	Down	0	0	AAEL012742-PA OS=Aedes aegypti GN=AAEL012742 PE=4 SV=1
CCG017643.1	1278	68	2675	2.28633956	73.17166695	5.000173539	Up	0	0	AAEL000294-PA OS=Aedes aegypti GN=AAEL000294 PE=4 SV=1
CCG009954.2	4675	467	2677	4.29237777	20.01781966	2.221435888	Up	5.22E-14	1.75E-13	AAEL008662-PA (Fragment) OS=Aedes aegypti GN=AAEL008662 PE=4 SV=1
CCG026216.1	2036	746	2678	15.7443134	45.98146621	1.546221629	Up	0	0	AAEL003943-PA OS=Aedes aegypti GN=AAEL003945 PE=4 SV=1
CCG028106.1	3319	975	2680	12.622926	28.22783421	1.161072081	Up	3.98E-12	9.20E-12	AAEL002643-PA OS=Aedes aegypti GN=AAEL002645 PE=4 SV=1
CCG022849.1	1548	516	2682	14.3232449	60.56724697	2.08017947	Up	9.75E-14	3.07E-13	AAEL005221-PA OS=Aedes aegypti GN=AAEL005221 PE=4 SV=1
CCG001161.1	4374	355	2683	3.48748418	21.44331426	2.620269329	Up	2.28E-13	6.52E-13	AAEL003768-PA OS=Aedes aegypti GN=AAEL003768 PE=4 SV=1
CCG025070.1	1773	11154	2690	270.323982	53.03878427	-2.349569894	Down	0	0	AAEL006586-PA (Fragment) OS=Aedes aegypti GN=AAEL006586 PE=3 SV=1
CCG011991.2	4995	530	2692	4.55935122	18.84037659	2.046927349	Up	0	0	AAEL012779-PA OS=Aedes aegypti GN=AAEL012779 PE=4 SV=1
CCG026098.1	26029	12320	2692	20.3383584	3.615493529	-2.491938737	Down	0	0	AAEL008102-PA OS=Aedes aegypti GN=AAEL008102 PE=4 SV=1
CCG017585.1	3171	209	2694	2.83212694	29.69965236	3.390488207	Up	0	0	AAEL001019-PA OS=Aedes aegypti GN=AAEL001019 PE=4 SV=1
CCG000682.1	1626	731	2694	19.3178819	57.91980174	1.584119743	Up	0	0	AAEL002178-PA (Fragment) OS=Aedes aegypti GN=AAEL002178 PE=4 SV=1
CCG021441.1	1717	949	2697	23.7497252	54.91116625	1.209188733	Up	1.30E-12	3.15E-12	AAEL004148-PA OS=Aedes aegypti GN=AAEL004148 PE=4 SV=1
CCG009109.3	3723	396	2701	4.57051166	25.36188707	2.472234508	Up	4.78E-13	1.24E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008434 PE=4 SV=1
CCG024743.2	3876	537	2701	5.95323723	24.36075995	2.03281285	Up	3.67E-13	9.84E-13	AAEL004950-PA OS=Aedes aegypti GN=AAEL004950 PE=4 SV=1

CCG002917.1	2730	1094	2701	17.2193735	34.58692511	1.006194106	Up	5.40E-12	1.22E-11	Mitochondrial Rho GTPase OS=Aedes aegypti GN=AAEL001853 PE=3 SV=1
CCG024591.1	1743	452	2702	11.1430408	54.19234873	2.281946201	Up	1.12E-13	3.49E-13	Uncharacterized protein OS=Anopheles darlingi GN=AND_07472 PE=4 SV=1
CCG003237.2	13052	958	2702	3.15392321	7.236995391	1.198243317	Up	1.10E-12	2.71E-12	AAEL004181-PA (Fragment) OS=Aedes aegypti GN=AAEL004181 PE=4 SV=1
CCG011828.1	1877	6137	2702	140.492947	50.32352895	-1.481192705	Down	0	0	AAEL010479-PA OS=Aedes aegypti GN=AAEL010479 PE=3 SV=1
CCG021145.1	693	26828	2705	1663.48058	136.4533026	-3.607725804	Down	0	0	AAEL011741-PA OS=Aedes aegypti GN=GSTS1 PE=3 SV=2
CCG006421.1	1212	0	2706	0.001	78.05041002	16.25211859	Up	4.44E-16	1.65E-15	AAEL004172-PA OS=Aedes aegypti GN=AAEL004172 PE=3 SV=1
CCG025409.1	931	424	2706	19.5694602	101.6080526	2.376338873	Up	0	0	60S acidic ribosomal protein P2 OS=Aedes aegypti GN=AAEL014583 PE=4 SV=1
CCG006087.2	1339	4	2707	0.12836366	70.67367828	9.104792376	Up	2.95E-05	5.03E-05	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG009479.1	982	7413	2707	324.373363	96.36665501	-1.75104942	Down	0	0	AAEL004987-PA OS=Aedes aegypti GN=AAEL004987 PE=4 SV=1
CCG014137.1	5497	936	2708	7.31665847	17.22157786	1.234960508	Up	1.10E-12	2.70E-12	AAEL003105-PA (Fragment) OS=Aedes aegypti GN=AAEL003105 PE=4 SV=1
CCG020879.1	358	15158	2708	1819.37217	264.4329986	-2.782466564	Down	0	0	AAEL000182-PA OS=Aedes aegypti GN=AAEL000182 PE=4 SV=1
CCG014784.1	3665	61	2709	0.71518521	25.83955574	5.175120544	Up	1.75E-14	6.17E-14	AAEL013142-PA OS=Aedes aegypti GN=AAEL013142 PE=4 SV=1
CCG021824.1	3001	304	2710	4.3528155	31.56845387	2.858462827	Up	0	0	AAEL005893-PA OS=Aedes aegypti GN=AAEL005893 PE=4 SV=1
CCG006572.1	2019	92	2715	1.95800673	47.00927263	4.58548773	Up	1.44E-14	5.12E-14	AAEL005744-PA OS=Aedes aegypti GN=AAEL005744 PE=4 SV=1
CCG025987.1	1776	284	2715	6.87128639	53.44128459	2.959302567	Up	0	0	AAEL008431-PA OS=Aedes aegypti GN=AAEL008431 PE=4 SV=1

CCG026566.1	3573	830	2716	9.98177434	26.57337803	1.412613442	Up	0	0	AAEL010074-PA OS=Aedes aegypti GN=AAEL010074 PE=4 SV=1
CCG025083.1	1479	11	2717	0.31958559	64.22017444	7.650680435	Up	2.67E-07	5.04E-07	AAEL001414-PA OS=Aedes aegypti GN=AAEL001414 PE=4 SV=1
CCG026597.1	703	282	2717	17.2367925	135.1090156	2.970560702	Up	5.45E-13	1.40E-12	-
CCG021430.1	6332	726	2718	4.92672573	15.00577957	1.606817207	Up	2.70E-13	7.54E-13	Target of rapamycin OS=Aedes aegypti GN=TOR PE=2 SV=1
CCG013245.1	645	16290	2719	1085.23562	147.3667512	-2.880525344	Down	0	0	L(2) long form OS=Culex quinquefasciatus GN=CpipJ_CPIJ017149 PE=4 SV=1
CCG015599.1	2902	123	2722	1.8212534	32.7899481	4.17025005	Up	0	0	AAEL010086-PA OS=Aedes aegypti GN=AAEL010086 PE=3 SV=1
CCG018192.1	1818	528	2722	12.4796589	52.34127028	2.068370436	Up	0	0	AGAP002194-PA OS=Anopheles gambiae GN=AgaP_AGAP002194 PE=4 SV=4
CCG001461.1	3022	384	2725	5.4600854	31.52260232	2.529391218	Up	1.03E-12	2.55E-12	AAEL010007-PA OS=Aedes aegypti GN=AAEL010007 PE=4 SV=1
CCG020336.1	3483	441	2727	5.44061239	27.37043375	2.330777343	Up	0	0	AAEL013686-PA OS=Aedes aegypti GN=AAEL013686 PE=4 SV=1
CCG025439.1	2500	6073	2728	104.382079	38.14647161	-1.452252521	Down	0	0	Brantneo-chain-amino-acid aminotransferase OS=Culex quinquefasciatus GN=CpipJ_CPIJ015408 PE=3 SV=1
CCG013806.1	1368	416	2732	13.0668199	69.81433636	2.417615254	Up	1.20E-14	4.28E-14	AAEL012312-PA OS=Aedes aegypti GN=AAEL012312 PE=4 SV=1
CCG002162.2	4332	8484	2732	84.1540232	22.04663254	-1.932473933	Down	0	0	AAEL005430-PA OS=Aedes aegypti GN=AAEL005430 PE=4 SV=1
CCG000169.1	2177	369	2733	7.28334041	43.88652752	2.591105942	Up	0	0	AAEL004920-PA OS=Aedes aegypti GN=AAEL004920 PE=4 SV=1
CCG016743.1	2421	512	2741	9.0873623	39.57894946	2.122799817	Up	3.04E-13	8.40E-13	AAEL002771-PA (Fragment) OS=Aedes aegypti GN=AAEL002771 PE=4 SV=1
CCG006489.1	968	10067	2742	446.876362	99.02437491	-2.174020139	Down	0	0	AAEL008789-PA OS=Aedes aegypti GN=AAEL008789 PE=4 SV=1

CCG028589.1	1104	698	2743	27.167459	86.85738513	1.676768884	Up	0	0	Eukaryotic translation initiation factor OS=Forcipomyia taiwana PE=2 SV=1
CCG000831.1	3918	952	2743	10.4408339	24.47436273	1.229034347	Up	4.77E-13	1.24E-12	Tripeptidyl-peptidase 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ013852 PE=4 SV=1
CCG013963.1	1882	93	2744	2.12337158	50.96998484	4.585219159	Up	4.04E-14	1.38E-13	AAEL003950-PA OS=Aedes aegypti GN=AAEL003950 PE=4 SV=1
CCG014817.2	3474	365	2745	4.51466699	27.62247258	2.613150984	Up	2.91E-13	8.05E-13	AAEL003162-PA OS=Aedes aegypti GN=AAEL003162 PE=4 SV=1
CCG013528.1	8169	565	2746	2.97195496	11.75118472	1.983322057	Up	4.06E-13	1.08E-12	Neurobeachin OS=Culex quinquefasciatus GN=CpipJ_CPIJ007145 PE=4 SV=1
CCG025566.1	2334	838	2746	15.4278653	41.12914654	1.41462268	Up	0	0	AAEL011432-PA OS=Aedes aegypti GN=AAEL011432 PE=4 SV=1
CCG013219.1	2108	875	2747	17.836109	45.55521171	1.352815192	Up	1.12E-12	2.74E-12	Methionine aminopeptidase OS=Aedes aegypti GN=AAEL003769 PE=3 SV=1
CCG021999.1	26232	804	2748	1.31700467	3.662143358	1.475427801	Up	2.72E-13	7.59E-13	Zinc finger protein 363 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000911 PE=4 SV=1
CCG026824.1	1611	4	2755	0.10669084	59.78277623	9.130149807	Up	2.95E-05	5.03E-05	AAEL000739-PA OS=Aedes aegypti GN=AAEL000739 PE=4 SV=1
CCG007750.2	4556	860	2755	8.11105614	21.13916868	1.381956958	Up	1.25E-12	3.05E-12	AAEL006054-PA OS=Aedes aegypti GN=AAEL006054 PE=4 SV=1
CCG014647.1	3537	591	2756	7.17984539	27.2391888	1.923659056	Up	0	0	AAEL004431-PA OS=Aedes aegypti GN=AAEL004431 PE=4 SV=1
CCG025390.1	2198	381	2758	7.44834799	43.8648441	2.558072758	Up	0	0	AAEL001520-PA OS=Aedes aegypti GN=AAEL001520 PE=4 SV=1
CCG017989.1	5540	780	2759	6.04989043	17.40972665	1.524912631	Up	1.45E-12	3.51E-12	AAEL001416-PA OS=Aedes aegypti GN=AAEL001416 PE=4 SV=1
CCG003940.1	4047	355	2760	3.76927497	23.8410783	2.661090541	Up	2.28E-13	6.52E-13	DNA repair protein RAD50 OS=Culex quinquefasciatus GN=CpipJ_CPIJ004115 PE=4 SV=1

CCG015130.1	7440	898	2760	5.18640076	12.968393	1.322194121	Up	2.22E-12	5.28E-12	AAEL011417-PA (Fragment) OS=Aedes aegypti GN=AAEL011417 PE=4 SV=1
CCG003530.1	1326	1018	2760	32.9888309	72.76383401	1.141243909	Up	4.26E-12	9.81E-12	AAEL009345-PA OS=Aedes aegypti GN=AAEL009345 PE=2 SV=1
CCG010112.1	1422	9361	2763	282.868977	67.9252593	-2.05811389	Down	0	0	AAEL005610-PA OS=Aedes aegypti GN=AAEL005610 PE=4 SV=1
CCG008786.1	2734	267	2764	4.19638593	35.34187162	3.074159173	Up	5.21E-13	1.35E-12	AAEL008519-PA OS=Aedes aegypti GN=AAEL008519 PE=4 SV=1
CCG028601.1	1071	196	2766	7.86374228	90.28440108	3.521188801	Up	7.93E-14	2.55E-13	AAEL001958-PA OS=Aedes aegypti GN=AAEL001958 PE=4 SV=1
CCG017093.1	1958	66	2767	1.44841802	49.40222259	5.092026013	Up	0	0	AAEL003355-PA OS=Aedes aegypti GN=AAEL003355 PE=4 SV=1
CCG018799.1	4383	582	2767	5.705769	22.06925663	1.95154479	Up	1.68E-13	4.94E-13	AAEL004039-PA OS=Aedes aegypti GN=AAEL004039 PE=4 SV=1
CCG004668.1	1059	20	2769	0.81151529	91.40648572	6.815534447	Up	3.69E-10	7.72E-10	AAEL013659-PA (Fragment) OS=Aedes aegypti GN=AAEL013659 PE=4 SV=1
CCG020863.1	1296	306	2769	10.1456318	74.69094783	2.880074699	Up	0	0	Histone deacetylase OS=Aedes aegypti GN=AAEL004586 PE=3 SV=1
CCG004765.2	519	44	2770	3.64290621	186.5788568	5.678551846	Up	3.91E-14	1.35E-13	AAEL009985-PA OS=Aedes aegypti GN=AAEL009985 PE=4 SV=1
CCG022083.1	2613	754	2770	12.3992269	37.05871667	1.579562751	Up	0	0	AGAP005609-PB OS=Anopheles gambiae GN=AGAP005609 PE=4 SV=4
CCG014857.1	1602	7793	2770	209.028178	60.44595921	-1.789979636	Down	0	0	-
CCG013224.2	876	1037	2773	50.8671402	110.6613031	1.121344926	Up	5.05E-12	1.15E-11	Cyclic-AMP response element binding protein OS=Aedes aegypti PE=2 SV=1
CCG008931.1	26148	1124	2774	1.84710042	3.70866834	1.005638956	Up	3.38E-12	7.88E-12	UDP-glucose 6-dehydrogenase OS=Aedes aegypti GN=AAEL011242 PE=3 SV=1
CCG023492.1	3061	683	2776	9.58782383	31.70342252	1.725363288	Up	1.46E-12	3.53E-12	AAEL004068-PA OS=Aedes aegypti GN=AAEL004068 PE=4 SV=1

CCG007455.1	743	9	2779	0.52049477	130.7524242	7.972738322	Up	1.09E-06	2.01E-06	-
CCG024097.1	2097	35	2779	0.7171868	46.32763526	6.013380306	Up	6.39E-14	2.10E-13	AAEL013822-PA OS=Aedes aegypti GN=AAEL013822 PE=4 SV=1
CCG023506.1	1119	10410	2782	399.745252	86.911462	-2.20146254	Down	0	0	AAEL002764-PA OS=Aedes aegypti GN=AAEL002764 PE=3 SV=1
CCG007160.1	2715	389	2783	6.15662128	35.83384319	2.541112052	Up	0	0	AAEL007558-PA (Fragment) OS=Aedes aegypti GN=AAEL007558 PE=4 SV=1
CCG018474.1	2091	436	2783	8.95973424	46.52744345	2.376554072	Up	1.61E-13	4.76E-13	AAEL007158-PA OS=Aedes aegypti GN=AAEL007158 PE=4 SV=1
CCG010643.1	3627	443	2783	5.2483023	26.82351372	2.353575509	Up	0	0	AAEL008382-PA OS=Aedes aegypti GN=AAEL008382 PE=4 SV=1
CCG016454.1	2673	1104	2783	17.7473202	36.39688899	1.03621394	Up	4.57E-12	1.05E-11	AGAP005609-PB OS=Anopheles gambiae GN=AGAP005609 PE=4 SV=4
CCG025990.1	1478	658	2787	19.129963	65.91929456	1.784866717	Up	1.34E-12	3.25E-12	AAEL010809-PA OS=Aedes aegypti GN=AAEL010809 PE=3 SV=1
CCG016818.1	3937	383	2789	4.18019008	24.76470254	2.566644841	Up	0	0	AAEL014695-PA OS=Aedes aegypti GN=AAEL014695 PE=4 SV=1
CCG005718.1	3892	4760	2790	52.5529128	25.06001855	-1.068383248	Down	6.34E-221	3.84E-220	-
CCG000378.1	1278	852	2793	28.6464897	76.39942647	1.415203442	Up	1.12E-12	2.74E-12	AAEL002160-PA OS=Aedes aegypti GN=AAEL002160 PE=4 SV=1
CCG003683.2	1555	666	2794	18.4037577	62.81249216	1.771051142	Up	0	0	AAEL009766-PA OS=Aedes aegypti GN=AAEL009766 PE=3 SV=1
CCG004256.1	1798	942	2796	22.5125083	54.3622591	1.2718786	Up	8.72E-13	2.18E-12	AAEL007556-PA OS=Aedes aegypti GN=AAEL007556 PE=4 SV=1
CCG025579.1	1838	540	2797	12.6244052	53.19820464	2.075162145	Up	9.26E-13	2.31E-12	AAEL011254-PA OS=Aedes aegypti GN=AAEL011254 PE=4 SV=1
CCG023435.1	3021	129	2799	1.83485461	32.3893468	4.14178172	Up	0	0	AAEL009411-PA OS=Aedes aegypti GN=AAEL009411 PE=4 SV=1
CCG019427.1	1953	158	2799	3.47630213	50.10149344	3.849228227	Up	0	0	AAEL003388-PA OS=Aedes aegypti GN=AAEL003388 PE=4 SV=1
CCG009524.1	3181	172	2800	2.32341853	30.77119615	3.727259561	Up	1.35E-13	4.09E-13	DNA polymerase OS=Aedes aegypti GN=AAEL009851 PE=3 SV=1

CCG026756.1	5127	398	2801	3.33566498	19.09852413	2.517414851	Up	7.64E-13	1.92E-12	AAEL004393-PA OS=Aedes aegypti GN=AAEL004393 PE=4 SV=1
CCG001810.1	3684	962	2804	11.2206527	26.60776549	1.245690754	Up	1.62E-12	3.89E-12	Structural maintenance of chromosomes protein OS=Aedes aegypti GN=AAEL005802 PE=3 SV=1
CCG026047.1	7072	364	2815	2.21167752	13.91509461	2.653437771	Up	0	0	Adenylyate cyclase type OS=Culex quinquefasciatus GN=CpipJ_CPIJ017287 PE=3 SV=1
CCG014305.1	3957	137	2816	1.48770625	24.87806606	4.063712739	Up	2.90E-13	8.04E-13	AAEL010566-PA OS=Aedes aegypti GN=AAEL010568 PE=3 SV=1
CCG014806.1	1596	603	2821	16.2348057	61.79028745	1.928289963	Up	3.84E-13	1.03E-12	AAEL013318-PA OS=Aedes aegypti GN=AAEL013318 PE=4 SV=1
CCG004957.1	2922	16492	2823	242.524594	33.77385877	-2.844152129	Down	0	0	AAEL000753-PA OS=Aedes aegypti GN=AAEL000753 PE=4 SV=1
CCG017985.2	4809	679	2827	6.06705132	20.55043636	1.760101607	Up	0	0	SYP1-dependent RNA helicase A OS=Culex quinquefasciatus GN=CpipJ_CPIJ003405 PE=4 SV=1
CCG012556.1	2412	135	2828	2.40502246	40.98756497	4.091064012	Up	1.90E-13	5.48E-13	AAEL011682-PA OS=Aedes aegypti GN=AAEL011682 PE=4 SV=1
CCG011175.1	1666	762	2832	19.6536241	59.42487384	1.596271566	Up	5.84E-13	1.50E-12	AAEL008366-PA OS=Aedes aegypti GN=AAEL008366 PE=4 SV=1
CCG026358.1	1461	6	2834	0.17646708	67.81092154	8.585974746	Up	8.38E-06	1.47E-05	AAEL015402-PA OS=Aedes aegypti GN=AAEL015402 PE=4 SV=1
CCG023968.1	5007	717	2840	6.1532454	19.82854125	1.68815911	Up	1.89E-14	6.63E-14	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001573 PE=4 SV=1
CCG017289.1	18896	693	2842	1.57589046	5.257801788	1.738292501	Up	1.99E-12	4.75E-12	AAEL012779-PA OS=Aedes aegypti GN=AAEL012779 PE=4 SV=1
CCG021231.1	6339	453	2843	3.07071932	15.67855827	2.352144347	Up	9.90E-13	2.45E-12	AAEL007242-PA (Fragment) OS=Aedes aegypti GN=AAEL007242 PE=4 SV=1
CCG005460.1	10959	7879	2847	30.8931964	9.08168756	-1.766256838	Down	0	0	Unm OS=Culex quinquefasciatus GN=CpipJ_CPIJ006896 PE=4 SV=1

CCG016447.1	2067	111	2848	2.3075184	48.1669919	4.383630768	Up	0	0	AAEL007271-PA OS=Aedes aegypti GN=AAEL007271 PE=4 SV=1
CCG006957.1	2034	128	2849	2.70409343	48.96564923	4.17855311	Up	4.15E-14	1.42E-13	AAEL005424-PA OS=Aedes aegypti GN=AAEL005424 PE=4 SV=1
CCG016148.1	1619	867	2849	23.0109697	61.51706641	1.418664927	Up	0	0	AAEL009601-PA OS=Aedes aegypti GN=AAEL009601 PE=4 SV=1
CCG022243.1	2698	918	2850	14.6205398	36.92775715	1.336709064	Up	1.42E-12	3.43E-12	AAEL012394-PA OS=Aedes aegypti GN=AAEL012394 PE=4 SV=1
CCG011837.1	43266	1029	2850	1.02195389	2.302757103	1.172032141	Up	3.54E-12	8.24E-12	AGAP001731-PA (Fragment) OS=Anopheles gambiae GN=AgaP_AGAP001731 PE=4 SV=5
CCG024324.1	1866	71	2851	1.63496847	53.41160079	5.029818408	Up	0	0	-
CCG019698.1	951	841	2851	37.9995235	104.8013113	1.463603538	Up	1.23E-12	3.01E-12	AAEL006820-PA OS=Aedes aegypti GN=AAEL006820 PE=4 SV=1
CCG022438.1	3942	266	2856	2.89953054	25.32745775	3.126811032	Up	4.09E-14	1.40E-13	AAEL007436-PA (Fragment) OS=Aedes aegypti GN=AAEL007436 PE=4 SV=1
CCG020990.1	3948	701	2857	7.6296312	25.29782086	1.729327891	Up	1.08E-12	2.64E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002576 PE=4 SV=1
CCG017626.1	2577	383	2861	6.38626634	38.81087693	2.603416404	Up	0	0	AAEL002446-PA (Fragment) OS=Aedes aegypti GN=AAEL002446 PE=4 SV=1
CCG012209.1	462	626	2862	58.2230603	216.5597146	1.895102314	Up	0	0	Superoxide dismutase [Cu-Zn] OS=Aedes aegypti GN=CUSOD3_b PE=2 SV=1
CCG010116.1	4563	338	2865	3.1829433	21.94947687	2.785753191	Up	1.44E-13	4.32E-13	AAEL005597-PA OS=Aedes aegypti GN=AAEL005597 PE=4 SV=1
CCG021521.1	1521	1038	2866	29.3245132	65.87141435	1.16754537	Up	2.33E-12	5.51E-12	AAEL005368-PA OS=Aedes aegypti GN=AAEL005368 PE=4 SV=1
CCG003220.2	1361	8796	2867	277.70888	73.64098421	-1.914992496	Down	0	0	AAEL009387-PA OS=Aedes aegypti GN=AAEL009387 PE=3 SV=1
CCG024704.1	5582	748	2869	5.75803681	17.96762738	1.641750997	Up	4.45E-13	1.17E-12	AAEL008615-PA OS=Aedes aegypti GN=AAEL008615 PE=4 SV=1

CCG016933.1	5968	923	2869	6.64562082	16.80551207	1.338458619	Up	0	0	AAEL012702-PA OS=Aedes aegypti GN=AAEL012702 PE=3 SV=1
CCG025244.1	2455	733	2872	12.8296601	40.89619996	1.67248385	Up	0	0	AAEL009755-PA OS=Aedes aegypti GN=AAEL009755 PE=4 SV=1
CCG004093.1	13503	911	2874	2.89901712	7.440575238	1.359850306	Up	0	0	AAEL011552-PA OS=Aedes aegypti GN=AAEL011552 PE=4 SV=1
CCG012284.1	3087	778	2875	10.829431	32.557514	1.5880331	Up	0	0	AAEL005261-PA OS=Aedes aegypti GN=AAEL005261 PE=4 SV=1
CCG003296.1	3364	700	2878	8.94138354	29.90782418	1.741952969	Up	0	0	AAEL006415-PA OS=Aedes aegypti GN=AAEL006415 PE=3 SV=1
CCG026678.1	2502	163	2879	2.79938719	40.22577091	3.844937123	Up	1.80E-13	5.24E-13	AAEL003950-PA OS=Aedes aegypti GN=AAEL003950 PE=4 SV=1
CCG000469.1	5895	946	2879	6.89556725	17.07292262	1.307968904	Up	0	0	AAEL014868-PA OS=Aedes aegypti GN=AAEL014868 PE=4 SV=1
CCG002462.1	2478	454	2886	7.87258253	40.71411895	2.370620301	Up	3.16E-13	8.66E-13	AAEL013968-PA OS=Aedes aegypti GN=AAEL013968 PE=4 SV=1
CCG016591.1	3680	506	2886	5.90833851	27.41564858	2.214175214	Up	3.38E-13	9.20E-13	Zinc finger protein ZTF1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000636 PE=4 SV=1
CCG014302.1	2922	1169	2888	17.190835	34.55150695	1.007109016	Up	2.14E-12	5.08E-12	AAEL006519-PA OS=Aedes aegypti GN=AAEL006519 PE=4 SV=1
CCG022878.1	3177	257	2889	3.47599049	31.78925452	3.193043143	Up	4.14E-13	1.09E-12	AAEL009912-PA OS=Aedes aegypti GN=AAEL009912 PE=4 SV=1
CCG019291.1	594	6313	2891	456.680025	170.1420508	-1.424443938	Down	0	0	AAEL007490-PA OS=Aedes aegypti GN=AAEL007490 PE=4 SV=1
CCG019772.1	4026	666	2892	7.10825714	25.11160865	1.820786674	Up	0	0	AAEL005495-PA (Fragment) OS=Aedes aegypti GN=AAEL005495 PE=4 SV=1
CCG005441.1	2962	816	2892	11.8377122	34.13211898	1.527739699	Up	2.12E-12	5.04E-12	Ubiquitin carboxyl-terminal hydrolase OS=Aedes aegypti GN=AAEL004980 PE=3 SV=1
CCG007200.2	5040	574	2894	4.89377533	20.07326448	2.036255484	Up	3.10E-13	8.52E-13	AAEL001470-PA OS=Aedes aegypti GN=AAEL001470 PE=4 SV=1
CCG013141.3	2034	239	2897	5.04904945	49.79062331	3.301790365	Up	6.72E-13	1.70E-12	AAEL004623-PA OS=Aedes aegypti GN=AAEL004623 PE=4 SV=1

CCG014852.1	2815	1007	2899	15.3714113	36.00143672	1.227804854	Up	3.64E-12	8.45E-12	AAEL006531-PA OS=Aedes aegypti GN=AAEL006531 PE=4 SV=1
CCG012278.1	1486	195	2901	5.6386933	68.24627249	3.59731747	Up	2.26E-13	6.47E-13	AAEL011199-PA OS=Aedes aegypti GN=AAEL011199 PE=4 SV=1
CCG019692.1	19554	811	2902	1.78216502	5.188141515	1.541586904	Up	0	0	Uncharacterized protein OS=Monodelphis domestica PE=4 SV=2
CCG021761.1	791	422	2906	22.9244349	128.4307867	2.486033003	Up	5.52E-13	1.42E-12	-
CCG012312.1	1614	431	2908	11.4745698	62.98554452	2.456580699	Up	0	0	AAEL007827-PA OS=Aedes aegypti GN=AAEL007827 PE=4 SV=1
CCG000668.1	980	668	2909	29.2895742	103.7690073	1.824916492	Up	9.28E-13	2.31E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_08855 PE=4 SV=1
CCG001357.1	3963	5415	2909	58.713377	25.66076889	-1.194124838	Down	2.92E-300	1.84E-299	Inhibitor of apoptosis protein 1 OS=Aedes albopictus GN=IAP1 PE=2 SV=1
CCG020320.1	1725	831	2911	20.7002026	58.99335866	1.510907661	Up	5.11E-15	1.84E-14	Zinc finger protein 442 OS=Culex quinquefasciatus GN=CpipJ_CPIJ011952 PE=4 SV=1
CCG006866.1	1119	273	2912	10.4832328	90.97274527	3.117350703	Up	1.53E-14	5.42E-14	AAEL009400-PA OS=Aedes aegypti GN=AAEL009400 PE=4 SV=1
CCG024680.1	1602	464	2912	12.445666	63.54463293	2.352126849	Up	0	0	AAEL006048-PA OS=Aedes aegypti GN=AAEL006048 PE=4 SV=1
CCG003322.1	2808	600	2914	9.18156722	36.27792682	1.982279675	Up	0	0	AGAP001361-PA OS=Anopheles gambiae GN=AgaP_AGAP001361 PE=4 SV=3
CCG010561.1	2313	589	2915	10.9421417	44.0567993	2.009469548	Up	0	0	AAEL012882-PA OS=Aedes aegypti GN=AAEL012882 PE=4 SV=1
CCG008069.1	2922	548	2916	8.05866344	34.88649386	2.114056125	Up	9.78E-13	2.42E-12	AAEL009605-PA OS=Aedes aegypti GN=AAEL009605 PE=4 SV=1
CCG017297.1	516	708	2918	58.9584731	197.6904101	1.745471822	Up	9.24E-13	2.30E-12	AAEL005369-PA OS=Aedes aegypti GN=AAEL005369 PE=4 SV=1
CCG028468.1	2862	5457	2918	81.9307623	35.64229616	-1.200814957	Down	2.00E-305	1.26E-304	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004708 PE=4 SV=1

CCG016744.1	1433	477	2924	14.3032543	71.33147333	2.318195344	Up	0	0	AAEL002771-PA (Fragment) OS=Aedes aegypti GN=AAEL002771 PE=4 SV=1
CCG021267.1	579	627	2931	46.5319924	176.9649555	1.927168824	Up	3.40E-13	9.23E-13	AAEL002705-PA OS=Aedes aegypti GN=AAEL002705 PE=4 SV=1
CCG010351.1	1717	695	2931	17.3931075	59.67542762	1.778621289	Up	1.60E-12	3.84E-12	AAEL003970-PA (Fragment) OS=Aedes aegypti GN=AAEL003970 PE=4 SV=1
CCG013793.1	1587	224	2933	6.06504131	64.60782972	3.413119638	Up	2.04E-13	5.88E-13	AAEL013390-PA OS=Aedes aegypti GN=AAEL013390 PE=4 SV=1
CCG016493.2	3238	913	2934	12.1159258	31.67621496	1.38649531	Up	0	0	AAEL006185-PA OS=Aedes aegypti GN=AAEL006185 PE=4 SV=1
CCG028181.1	4957	982	2934	8.51246306	20.6914634	1.281387145	Up	2.51E-12	5.91E-12	AAEL006073-PA OS=Aedes aegypti GN=AAEL006073 PE=4 SV=1
CCG006303.1	3207	642	2936	8.60198616	32.00420973	1.89551997	Up	0	0	Testis-specific zinc finger protein topi OS=Culex quinquefasciatus GN=CpipJ_CPIJ001933 PE=4 SV=1
CCG005122.1	1715	429	2937	10.7486975	59.86732296	2.477606917	Up	0	0	AAEL003998-PA OS=Aedes aegypti GN=AAEL003998 PE=4 SV=1
CCG018015.1	5073	517	2939	4.37913519	20.25278443	2.209402377	Up	8.39E-14	2.69E-13	AAEL001384-PA (Fragment) OS=Aedes aegypti GN=AAEL001384 PE=4 SV=1
CCG025177.1	2500	942	2939	16.190996	41.09695017	1.343839598	Up	8.72E-13	2.18E-12	Glycylpeptide N- tetradecanoyltransferase OS=Aedes aegypti GN=AAEL005092 PE=3 SV=1
CCG019549.1	582	941	2940	69.4751207	176.5933569	1.345862731	Up	0	0	AAEL005038-PA OS=Aedes aegypti GN=AAEL005038 PE=3 SV=1
CCG014513.1	3081	989	2943	13.7932709	33.39247275	1.27555832	Up	5.21E-12	1.18E-11	GH16388 OS=Drosophila grimshawi GN=GH16388 PE=4 SV=1
CCG003475.2	2705	187	2944	2.97055097	38.04701176	3.6789807	Up	3.31E-13	9.04E-13	AAEL012224-PA OS=Aedes aegypti GN=AAEL012224 PE=4 SV=1

CCG004210.1	41633	7222	2944	7.4538809	2.472009387	-1.592307545	Down	0	0	AAEL008340-PA (Fragment) OS=Aedes aegypti GN=AAEL008340 PE=4 SV=1
CCG023651.1	1798	228	2946	5.44888737	57.27868931	3.393964905	Up	1.05E-13	3.28E-13	GK25828 OS=Drosophila willistoni GN=GK25828 PE=4 SV=1
CCG010879.1	3714	349	2952	4.03781297	27.78590012	2.782706984	Up	4.55E-14	1.54E-13	AAEL011287-PA OS=Aedes aegypti GN=AAEL011287 PE=4 SV=1
CCG024822.1	54217	9904	2954	7.84942456	1.90469317	-2.043028285	Down	0	0	GE15410 OS=Drosophila yakuba GN=GE15410 PE=4 SV=1
CCG026796.1	4251	10748	2954	108.642368	24.29234288	-2.161013291	Down	0	0	AAEL003837-PA OS=Aedes aegypti GN=AAEL003837 PE=4 SV=1
CCG000833.1	763	7897	2958	444.733937	135.5263207	-1.714369434	Down	0	0	AAEL009078-PA OS=Aedes aegypti GN=AAEL009078 PE=4 SV=1
CCG007976.1	2612	526	2961	8.65317014	39.62919507	2.19526299	Up	1.08E-13	3.37E-13	AAEL010457-PA OS=Aedes aegypti GN=AAEL010457 PE=4 SV=1
CCG006513.1	3438	61	2966	0.76240658	30.15888566	5.305878749	Up	1.75E-14	6.17E-14	AAEL009238-PA OS=Aedes aegypti GN=AAEL009238 PE=4 SV=1
CCG010024.1	1840	76	2967	1.77483686	56.3702213	4.989174902	Up	0	0	AAEL003382-PA OS=Aedes aegypti GN=AAEL003382 PE=4 SV=1
CCG021458.1	39389	888	2967	0.96872539	2.633253121	1.442686549	Up	0	0	AAEL004918-PA OS=Aedes aegypti GN=AAEL004918 PE=4 SV=1
CCG008993.1	423	320	2974	32.506655	245.7823053	2.918574041	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005950 PE=4 SV=1
CCG018200.1	1952	308	2976	6.78006058	53.29704492	2.974685474	Up	0	0	AAEL008630-PA OS=Aedes aegypti GN=AAEL008630 PE=4 SV=1
CCG021501.1	615	9	2977	0.62882538	169.2207967	8.072031709	Up	1.09E-06	2.01E-06	Putative salivary secreted 30 kDa allergen-like protein 30k-5 OS=Aedes albopictus PE=2 SV=1
CCG020676.1	2214	45	2979	0.87336859	47.03735614	5.751072516	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001409 PE=4 SV=1
CCG023405.1	660	305	2984	19.8572258	158.0537847	2.992679592	Up	5.30E-13	1.37E-12	AAEL010954-PA OS=Aedes aegypti GN=AAEL010954 PE=4 SV=1
CCG015183.1	309	0	2985	0.001	337.7037416	18.36539864	Up	4.44E-16	1.65E-15	-

CCG018630.1	2364	526	2985	9.56094772	44.14147892	2.206909431	Up	1.08E-13	3.37E-13	AAEL008677-FA OS=Aedes aegypti GN=AAEL008677 PE=4 SV=1
CCG005745.2	3702	292	2986	3.38929295	28.19703253	3.056487095	Up	0	0	AAEL006680-FA OS=Aedes aegypti GN=AAEL006680 PE=4 SV=1
CCG025505.1	20730	892	2987	1.84896301	5.037162215	1.445894826	Up	2.08E-13	5.98E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005572 PE=4 SV=1
CCG015107.1	1077	353	2992	14.0838592	97.11714401	2.78568329	Up	0	0	AAEL003179-FA OS=Aedes aegypti GN=AAEL003179 PE=4 SV=1
CCG013193.1	4005	1157	2992	12.4134789	26.11614584	1.073034515	Up	4.14E-12	9.55E-12	AAEL010871-FA OS=Aedes aegypti GN=AAEL010871 PE=4 SV=1
CCG021368.1	3663	15291	2993	179.374887	28.56405197	-2.650705371	Down	0	0	AAEL005422-FA OS=Aedes aegypti GN=AAEL005422 PE=3 SV=1
CCG027749.1	1791	494	2995	11.8520653	58.45898321	2.30228626	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003817 PE=4 SV=1
CCG023058.2	1314	161	2996	5.26493704	79.7069994	3.920218234	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012003 PE=4 SV=1
CCG020313.2	5085	1163	2996	9.82768955	20.59685294	1.067499731	Up	4.90E-12	1.12E-11	Multidrug resistance-associated protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ011963 PE=3 SV=1
CCG019627.1	48622	5233	2996	4.62466828	2.154066003	-1.102287431	Down	2.74E-255	1.69E-254	Monocarboxylate transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ014208 PE=4 SV=1
CCG027727.1	2790	109	3000	1.6787459	37.58954492	4.484875664	Up	0	0	AAEL013258-FA OS=Aedes aegypti GN=AAEL013258 PE=4 SV=1
CCG004218.1	2692	616	3004	9.832599	39.00990469	1.988195761	Up	3.02E-13	8.35E-13	AAEL005214-FA OS=Aedes aegypti GN=AAEL005214 PE=4 SV=1
CCG014517.2	1531	595	3008	16.6995376	68.6835379	2.040156197	Up	7.20E-13	1.81E-12	AAEL013484-FA OS=Aedes aegypti GN=AAEL013484 PE=4 SV=1
CCG003676.1	5801	902	3012	6.68138263	18.15106527	1.441835635	Up	3.95E-13	1.05E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007358 PE=4 SV=1

CCG020581.1	2676	917	3014	14.7246811	39.37378408	1.418998982	Up	0	0	AAEL013138-PA OS=Aedes aegypti GN=AAEL013138 PE=4 SV=1
CCG023745.1	3030	986	3014	13.9828905	34.77367861	1.314333069	Up	1.31E-12	3.18E-12	AAEL009264-PA OS=Aedes aegypti GN=AAEL009264 PE=4 SV=1
CCG028183.1	6465	571	3016	3.7951614	16.30845518	2.103386982	Up	0	0	Transcription initiation factor TFIID subunit 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003658 PE=4 SV=1
CCG011962.1	3453	385	3018	4.79100719	30.55432357	2.672975659	Up	7.93E-13	1.99E-12	AAEL008617-PA OS=Aedes aegypti GN=AAEL008617 PE=4 SV=1
CCG008911.1	1131	72144	3018	2740.94477	93.28388974	-4.8769015	Down	0	0	AAEL005961-PA OS=Aedes aegypti GN=AAEL005961 PE=3 SV=1
CCG019084.1	1080	715	3019	28.4475558	97.72133109	1.780368814	Up	0	0	AAEL005493-PA OS=Aedes aegypti GN=AAEL005493 PE=3 SV=1
CCG024930.1	2387	894	3024	16.0933987	44.28731837	1.460424607	Up	5.67E-13	1.46E-12	AAEL006155-PA OS=Aedes aegypti GN=AAEL006155 PE=4 SV=1
CCG028651.1	3999	1046	3028	11.2393954	26.47003302	1.235795558	Up	3.46E-12	8.05E-12	AAEL009678-PA OS=Aedes aegypti GN=AAEL009678 PE=4 SV=1
CCG017657.1	1959	647	3029	14.1916377	54.05238404	1.929317165	Up	0	0	AAEL000825-PA OS=Aedes aegypti GN=AAEL000825 PE=4 SV=1
CCG010157.1	5163	938	3029	7.8066262	20.50912654	1.393494954	Up	1.14E-12	2.79E-12	AAEL013012-PA (Fragment) OS=Aedes aegypti GN=AAEL013012 PE=4 SV=1
CCG019745.1	4620	270	3030	2.51121826	22.92718152	3.190599685	Up	2.59E-13	7.26E-13	AAEL012057-PA OS=Aedes aegypti GN=AAEL012057 PE=4 SV=1
CCG027752.1	6096	498	3032	3.51032281	17.38738438	2.30836531	Up	3.42E-13	9.27E-13	AAEL008557-PA (Fragment) OS=Aedes aegypti GN=AAEL008557 PE=4 SV=1
CCG014772.1	411	246	3035	25.7191112	258.1468857	3.3272795	Up	2.51E-13	7.05E-13	Histone H2A OS=Aedes aegypti GN=AAEL012499 PE=3 SV=1
CCG007979.1	1483	5558	3035	161.042336	71.54306811	-1.170556114	Down	1.68E-298	1.06E-297	AAEL004985-PA OS=Aedes aegypti GN=AAEL004985 PE=2 SV=1
CCG027289.1	2387	291	3036	5.23845529	44.4630617	3.085393936	Up	6.00E-14	2.00E-13	AAEL000978-PA (Fragment) OS=Aedes aegypti GN=AAEL000978 PE=4 SV=1

CCG023864.1	1554	12647	3036	349.702853	68.29686505	-2.356238304	Down	0	0	AAEL001094-PA OS=Aedes aegypti GN=AAEL001094 PE=3 SV=1
CCG028495.1	2483	499	3039	8.63548029	42.78622759	2.308798158	Up	0	0	AAEL0008530-PA OS=Aedes aegypti GN=AAEL008530 PE=4 SV=1
CCG016754.1	1443	6345	3039	188.941764	73.62314838	-1.359710285	Down	0	0	AAEL010479-PA OS=Aedes aegypti GN=AAEL010479 PE=3 SV=1
CCG002610.2	552	273	3042	21.2513361	192.6505035	3.180360501	Up	1.53E-14	5.41E-14	AAEL0006229-PA OS=Aedes aegypti GN=AAEL006229 PE=4 SV=1
CCG007043.1	6744	270	3046	1.72031856	15.789281	3.198197833	Up	2.59E-13	7.26E-13	AAEL001612-PA OS=Aedes aegypti GN=AAEL001612 PE=4 SV=1
CCG024526.1	5196	1108	3046	9.16290723	20.49324693	1.161271264	Up	4.45E-12	1.02E-11	AAEL005801-PA OS=Aedes aegypti GN=AAEL005801 PE=3 SV=1
CCG017744.1	2298	257	3047	4.80557955	46.35242355	3.269862439	Up	4.14E-13	1.10E-12	AAEL010847-PA OS=Aedes aegypti GN=AAEL010847 PE=4 SV=1
CCG012311.1	2460	576	3047	10.0612062	43.29994688	2.105561987	Up	7.26E-13	1.83E-12	AAEL007827-PA OS=Aedes aegypti GN=AAEL007827 PE=4 SV=1
CCG028182.1	3861	770	3047	8.56946274	27.58815574	1.686772353	Up	0	0	AAEL0006087-PA OS=Aedes aegypti GN=AAEL006087 PE=4 SV=1
CCG012479.1	1295	957	3047	31.754468	82.25318095	1.373111874	Up	1.76E-12	4.21E-12	Predicted ER protein with RDEL retention signal OS=Aedes aegypti PE=2 SV=1
CCG013429.1	2376	429	3050	7.7584243	44.87489232	2.532072894	Up	0	0	AAEL004068-PA OS=Aedes aegypti GN=AAEL004068 PE=4 SV=1
CCG017167.1	657	75	3051	4.90522085	162.3404908	5.048560978	Up	0	0	AAEL000695-PA OS=Aedes aegypti GN=AAEL000695 PE=4 SV=1
CCG022143.1	1701	11333	3054	286.288067	62.76459569	-2.189444642	Down	0	0	AAEL004034-PA OS=Aedes aegypti GN=CYP4G36 PE=3 SV=1
CCG018320.1	1562	138	3055	3.79630178	68.37230188	4.170745412	Up	1.49E-13	4.45E-13	AAEL003664-PA OS=Aedes aegypti GN=AAEL003664 PE=4 SV=1
CCG010685.1	1446	55	3056	1.63439516	73.88139268	5.498382318	Up	0	0	AAEL011653-PA OS=Aedes aegypti GN=AAEL011653 PE=3 SV=1
CCG020372.1	2874	738	3057	11.033982	37.18422133	1.752737035	Up	0	0	AAEL014329-PA (Fragment) OS=Aedes aegypti GN=AAEL014329 PE=4 SV=1

CCG012473.1	2124	475	3058	9.60952163	50.33070168	2.388902192	Up	6.44E-13	1.64E-12	AAEL005215-PA OS=Aedes aegypti GN=AAEL005215 PE=4 SV=1
CCG017119.1	3756	622	3058	7.11586127	28.4617706	1.999915125	Up	0	0	AAEL009096-PA OS=Aedes aegypti GN=AAEL009096 PE=4 SV=1
CCG004563.1	3191	1098	3059	14.7855746	33.51218071	1.180495256	Up	3.40E-12	7.92E-12	AAEL012286-PA OS=Aedes aegypti GN=AAEL012286 PE=4 SV=1
CCG010329.2	3948	250	3061	2.72098117	27.10417558	3.316316249	Up	2.10E-13	6.02E-13	AAEL000984-PA (Fragment) OS=Aedes aegypti GN=AAEL000984 PE=4 SV=1
CCG019746.1	3423	440	3061	5.52342484	31.26125773	2.50074082	Up	0	0	AAEL012057-PA OS=Aedes aegypti GN=AAEL012057 PE=4 SV=1
CCG025420.2	2643	698	3064	11.3480419	40.52673478	1.83643056	Up	0	0	AAEL009470-PA (Fragment) OS=Aedes aegypti GN=AAEL009470 PE=4 SV=1
CCG024597.1	2956	945	3068	13.7369415	36.28281229	1.401225452	Up	2.91E-12	6.82E-12	AAEL005522-PA OS=Aedes aegypti GN=AAEL005522 PE=4 SV=1
CCG020873.1	462	6999	3068	650.963577	232.1471713	-1.487537121	Down	0	0	AAEL000179-PA OS=Aedes aegypti GN=AAEL000179 PE=3 SV=1
CCG022630.1	1538	418	3076	11.6783804	69.91655354	2.58179386	Up	6.74E-13	1.70E-12	AAEL009309-PA OS=Aedes aegypti GN=AAEL009309 PE=4 SV=1
CCG003165.1	1692	287	3078	7.28860156	63.594312	3.125183793	Up	2.87E-13	7.99E-13	AAEL004317-PA OS=Aedes aegypti GN=AAEL004317 PE=4 SV=1
CCG004300.1	657	7430	3078	485.943878	163.7771323	-1.569055775	Down	0	0	AAEL003118-PA OS=Aedes aegypti GN=AAEL003118 PE=4 SV=1
CCG014526.1	3105	374	3079	5.17574259	34.66555046	2.743664896	Up	1.37E-13	4.12E-13	AAEL010827-PA OS=Aedes aegypti GN=AAEL010827 PE=4 SV=1
CCG012264.1	2163	743	3079	14.7602926	49.76261405	1.753340955	Up	0	0	AAEL010903-PA OS=Aedes aegypti GN=AAEL010903 PE=4 SV=1
CCG000540.1	2062	116	3080	2.41730806	52.21701865	4.433046844	Up	1.16E-13	3.59E-13	AAEL002843-PA OS=Aedes aegypti GN=AAEL002843 PE=4 SV=1
CCG017503.1	5069	367	3084	3.11104608	21.26875628	2.773264001	Up	4.12E-13	1.09E-12	AAEL009067-PA OS=Aedes aegypti GN=AAEL009067 PE=4 SV=1
CCG018786.1	1975	559	3087	12.1620667	54.64111412	2.167598499	Up	3.87E-13	1.03E-12	AAEL009647-PA OS=Aedes aegypti GN=AAEL009647 PE=4 SV=1
CCG011248.1	1711	1137	3087	28.5544058	63.07200491	1.143286433	Up	4.86E-12	1.11E-11	-

CCG022850.1	2259	546	3092	10.3857791	47.84904461	2.203880667	Up	3.03E-13	8.37E-13	AAEL005214-PA OS=Aedes aegypti GN=AAEL005214 PE=4 SV=1
CCG005892.1	4703	907	3093	8.28695498	22.99084628	1.472145581	Up	0	0	AAEL004942-PA OS=Aedes aegypti GN=AAEL004942 PE=4 SV=1
CCG018089.1	669	512	3096	32.8856564	161.7800073	2.29850296	Up	3.04E-13	8.39E-13	AAEL007947-PA OS=Aedes aegypti GN=GSTE3 PE=3 SV=1
CCG003444.1	751	723	3099	41.3676673	144.2552593	1.802048406	Up	0	0	AAEL012856-PA OS=Aedes aegypti GN=AAEL012856 PE=4 SV=1
CCG006156.2	3559	513	3101	6.19372685	30.45957187	2.298015999	Up	0	0	AAEL001440-PA OS=Aedes aegypti GN=AAEL001440 PE=4 SV=1
CCG012083.1	7665	736	3101	4.12599148	14.14293754	1.777269058	Up	0	0	AAEL006013-PA OS=Aedes aegypti GN=AAEL006013 PE=4 SV=1
CCG007418.1	4828	476	3103	4.23645271	22.46800597	2.406943422	Up	4.62E-13	1.21E-12	SV1 dependent RNA helicase A OS=Culex quinquefasciatus GN=CpipJ_CPIJ003405 PE=4 SV=1
CCG016957.1	1487	290	3105	8.38010964	72.99626723	3.122781667	Up	1.62E-13	4.77E-13	AAEL004484-PA OS=Aedes aegypti GN=AAEL004484 PE=4 SV=1
CCG007992.1	738	416	3106	24.2214222	147.1279237	2.6027156	Up	1.20E-14	4.28E-14	AAEL011314-PA OS=Aedes aegypti GN=AAEL011314 PE=2 SV=1
CCG013839.1	4026	681	3108	7.26835302	26.98716448	1.892573004	Up	1.31E-14	4.67E-14	AAEL000747-PA OS=Aedes aegypti GN=AAEL000747 PE=4 SV=1
CCG011968.1	1165	1069	3108	39.4288809	93.26208087	1.242037855	Up	3.42E-12	7.98E-12	Glutathione S-transferase OS=Aedes aegypti PE=2 SV=1
CCG026271.1	1719	101	3114	2.52469063	63.32755897	4.64865495	Up	0	0	AAEL002968-PA OS=Aedes aegypti GN=AAEL002968 PE=4 SV=1
CCG003864.1	4842	652	3114	5.78609396	22.48246053	1.958138279	Up	1.50E-12	3.61E-12	AAEL002455-PA OS=Aedes aegypti GN=AAEL002455 PE=4 SV=1
CCG000409.1	4404	663	3115	6.46887694	24.7263924	1.934464592	Up	1.08E-13	3.38E-13	AAEL007563-PA (Fragment) OS=Aedes aegypti GN=DUOX PE=4 SV=2
CCG024287.1	3142	773	3119	10.5714847	34.70237596	1.714856437	Up	5.30E-13	1.37E-12	AAEL003559-PA (Fragment) OS=Aedes aegypti GN=AAEL003559 PE=4 SV=1
CCG000194.1	3654	934	3119	10.9835063	29.8398646	1.441902301	Up	3.46E-13	9.34E-13	AAEL006145-PA OS=Aedes aegypti GN=AAEL006145 PE=4 SV=1

CCG006486.1	2935	6280	3121	91.9420557	37.17369056	-1.306442998	Down	0	0	ATP synthase alpha subunit vacuolar OS=Culex quinquefasciatus GN=CpipJ_CPIJ007772 PE=3 SV=1
CCG016531.1	2276	981	3122	18.5207863	47.95243413	1.3724587	Up	4.12E-12	9.52E-12	AAEL0006497-PA OS=Aedes aegypti GN=AAEL008497 PE=4 SV=1
CCG004947.1	5726	1155	3123	8.66748926	19.06648592	1.137352922	Up	4.50E-12	1.04E-11	AAEL001554-PA OS=Aedes aegypti GN=AAEL001554 PE=4 SV=1
CCG011505.1	6546	595	3124	3.90574276	16.68341837	2.094746084	Up	7.20E-13	1.81E-12	AAEL007857-PA OS=Aedes aegypti GN=AAEL007857 PE=4 SV=1
CCG005834.1	4822	576	3129	5.13284262	22.68445625	2.143874146	Up	7.26E-13	1.83E-12	AAEL011963-PA OS=Aedes aegypti GN=AAEL011963 PE=4 SV=1
CCG020875.1	1467	320	3131	9.37308458	74.6110188	2.992792902	Up	0	0	AAEL000193-PA OS=Aedes aegypti GN=AAEL000193 PE=4 SV=1
CCG002140.1	4272	494	3131	4.96887849	25.62134002	2.366353765	Up	0	0	AAEL010414-PA OS=Aedes aegypti GN=AAEL010414 PE=4 SV=1
CCG003695.1	1591	502	3131	13.5580181	68.79595511	2.343177443	Up	6.14E-13	1.57E-12	AAEL005064-PA OS=Aedes aegypti GN=CLIPB5 PE=3 SV=1
CCG004682.1	1326	1081	3132	35.0303794	82.57113337	1.237030893	Up	3.20E-12	7.49E-12	AAEL001810-PA OS=Aedes aegypti GN=AAEL001810 PE=4 SV=1
CCG022797.1	3516	698	3137	8.53039669	31.19002111	1.870399791	Up	0	0	SV=1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000640 PE=4 SV=1
CCG000616.1	2532	1157	3138	19.6350644	43.32506813	1.141769692	Up	4.14E-12	9.55E-12	Splicing factor 5B subunit 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ010596 PE=4 SV=1
CCG002298.2	2637	490	3139	7.98451648	41.61320849	2.381764578	Up	0	0	AAEL002145-PA OS=Aedes aegypti GN=AAEL002145 PE=4 SV=1
CCG016541.1	4536	606	3145	5.7406656	24.23804684	2.077983522	Up	3.75E-13	1.00E-12	AAEL007521-PA OS=Aedes aegypti GN=AAEL007521 PE=4 SV=1
CCG009903.1	3898	889	3147	9.79992152	28.22311365	1.526035058	Up	0	0	AAEL009040-PA OS=Aedes aegypti GN=AAEL009040 PE=4 SV=1
CCG022447.1	6822	198	3155	1.24714269	16.16730625	3.696380874	Up	0	0	AAEL010499-PA (Fragment) OS=Aedes aegypti GN=AAEL010499 PE=4 SV=1

CCG012959.1	5844	601	3156	4.41902986	18.87890512	2.094973513	Up	0	0	AAEL000386-PA OS=Aedes aegypti GN=AAEL000386 PE=4 SV=1
CCG025583.1	5250	793	3160	6.4904761	21.04155326	1.696844991	Up	0	0	AAEL000409-PA OS=Culex quinquefasciatus GN=CpipJ_CPIJ005144 PE=4 SV=1
CCG014949.1	26271	25961	3160	42.4626881	4.204946694	-3.336036289	Down	0	0	AAEL009863-PA OS=Aedes aegypti GN=AAEL009863 PE=4 SV=1
CCG024099.2	5318	656	3162	5.30051634	20.78564708	1.971382852	Up	9.98E-13	2.47E-12	AAEL007252-PA OS=Aedes aegypti GN=AAEL007252 PE=4 SV=1
CCG010173.1	3018	405	3163	5.76631627	36.63784938	2.667612947	Up	2.34E-13	6.64E-13	AAEL000409-PA OS=Aedes aegypti GN=AAEL006469 PE=4 SV=1
CCG019499.2	4642	225	3168	2.08276396	23.8577813	3.517888633	Up	0	0	AAEL009397-PA OS=Aedes aegypti GN=AAEL009397 PE=4 SV=1
CCG010270.1	6456	949	3168	6.316338	17.15424734	1.441405547	Up	1.30E-12	3.15E-12	AAEL004720-PA (Fragment) OS=Aedes aegypti GN=AAEL004720 PE=4 SV=1
CCG022003.2	1032	22	3173	0.91602147	107.4831514	6.874513391	Up	8.38E-11	1.80E-10	AAEL007097-PA OS=Aedes aegypti GN=AAEL007097 PE=4 SV=1
CCG010445.1	2766	1146	3174	17.8030788	40.11481217	1.172008288	Up	4.08E-12	9.43E-12	AAEL007470-PA OS=Aedes aegypti GN=AAEL007470 PE=4 SV=1
CCG021366.1	4482	317	3176	3.03913563	24.77186234	3.026969371	Up	2.22E-13	6.35E-13	AAEL003162-PA OS=Aedes aegypti GN=AAEL003162 PE=4 SV=1
CCG000065.1	654	1134	3179	74.5071545	169.9271588	1.18946558	Up	2.06E-12	4.92E-12	-
CCG021499.1	633	0	3183	0.001	175.7854581	17.4234562	Up	4.44E-16	1.63E-15	30 kDa salivary gland allergen OS=Aedes albopictus PE=2 SV=1
CCG010321.1	1909	756	3187	17.0168252	58.36146049	1.778054083	Up	0	0	AAEL000577-PA OS=Aedes aegypti GN=AAEL006577 PE=3 SV=1
CCG016391.1	2895	6902	3191	102.444597	38.53259454	-1.410692661	Down	0	0	AAEL013800-PA OS=Aedes aegypti GN=AAEL013800 PE=4 SV=1
CCG016660.1	6588	793	3192	5.17228287	16.93789002	1.711381084	Up	0	0	AAEL004375-PA OS=Aedes aegypti GN=AAEL004375 PE=4 SV=1
CCG018707.1	948	5676	3192	257.274487	117.7076155	-1.128100734	Down	2.94E-287	1.84E-286	AAEL003596-PA OS=Aedes aegypti GN=AAEL003596 PE=4 SV=1
CCG005064.1	1329	271	3193	8.7620753	83.98929852	3.260861	Up	0	0	AAEL005904-PA OS=Aedes aegypti GN=AAEL005904 PE=4 SV=1

CCG011099.1	7304	8427	3196	49.5763901	15.29663918	-1.696438513	Down	0	0	-
CCG004333.1	504	986	3199	84.0638062	221.8879512	1.400274645	Up	1.31E-12	3.18E-12	AGAP002251-PA OS=Anopheles gambiae GN=AGAP002251 PE=3 SV=4
CCG019681.1	1767	699	3200	16.9982142	63.30870723	1.897020748	Up	0	0	AAEL012750-PA OS=Aedes aegypti GN=AAEL012750 PE=4 SV=1
CCG015283.1	2414	5263	3204	93.6825656	46.39864075	-1.013698043	Down	6.18E-224	3.74E-223	AAEL000339-PA OS=Aedes aegypti GN=AAEL000339 PE=4 SV=1
CCG024423.1	2513	566	3206	9.6780222	44.59858151	2.204213671	Up	6.91E-14	2.25E-13	AAEL003539-PA OS=Aedes aegypti GN=AAEL003539 PE=4 SV=1
CCG022184.1	4899	1178	3206	10.3323836	22.87736994	1.14674809	Up	4.91E-12	1.12E-11	Rho GTPase-activating protein 5 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009693 PE=4 SV=1
CCG026969.1	4064	5364	3207	56.7149745	27.58641575	-1.03977168	Down	6.61E-238	4.04E-237	AAEL008723-PA OS=Aedes aegypti GN=AAEL008723 PE=4 SV=1
CCG002780.1	1641	583	3208	15.2659082	68.34012912	2.162419557	Up	2.70E-13	7.53E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000417 PE=4 SV=1
CCG001509.1	9205	774	3209	3.61309881	12.18697557	1.754031522	Up	0	0	DNA-directed RNA polymerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007655 PE=3 SV=1
CCG010776.1	4954	912	3212	7.91045579	22.66572164	1.518679367	Up	0	0	AAEL012935-PA OS=Aedes aegypti GN=AAEL012935 PE=4 SV=1
CCG000523.1	1468	1244	3213	36.4130449	76.51290413	1.071247699	Up	1.32E-12	3.20E-12	AAEL008645-PA OS=Aedes aegypti GN=AAEL008645 PE=4 SV=1
CCG022780.1	2466	303	3215	5.27973625	45.57617998	3.109742243	Up	1.37E-13	4.14E-13	AAEL011023-PA OS=Aedes aegypti GN=AAEL011023 PE=4 SV=1
CCG018879.1	4428	1051	3215	10.1990043	25.38185633	1.315369272	Up	3.63E-12	8.44E-12	AAEL012701-PA OS=Aedes aegypti GN=AAEL012701 PE=3 SV=1
CCG013119.1	6576	1166	3216	7.61902533	17.09638353	1.166012822	Up	2.46E-12	5.80E-12	ADAM metalloprotease OS=Culex quinquefasciatus GN=CpipJ_CPIJ012681 PE=4 SV=1
CCG026626.1	6468	1119	3218	7.43400325	17.39266151	1.226267494	Up	5.84E-12	1.32E-11	AAEL009219-PA OS=Aedes aegypti GN=AAEL009219 PE=4 SV=1

CCG001776.1	4818	306	3221	2.72908651	23.37081974	3.098218307	Up	0	0	AAEL005814-PA OS=Aedes aegypti GN=AAEL005814 PE=4 SV=1
CCG023664.2	5982	6588	3221	47.3227368	18.82323796	-1.330018691	Down	0	0	AAEL005858-PA OS=Aedes aegypti GN=AAEL003658 PE=4 SV=1
CCG011932.1	333	22227	3221	2868.13301	338.1399684	-3.08441947	Down	0	0	AAEL005234-PA OS=Aedes aegypti GN=AAEL003234 PE=3 SV=1
CCG026097.1	1894	818	3224	18.5582064	59.50659151	1.680992199	Up	0	0	AAEL008100-PA OS=Aedes aegypti GN=AAEL008100 PE=4 SV=1
CCG023865.1	3794	336	3226	3.80543775	29.72467076	2.965526504	Up	1.75E-13	5.12E-13	AAEL001099-PA OS=Aedes aegypti GN=AAEL001099 PE=4 SV=1
CCG012671.1	1617	772	3232	20.5149259	69.87331511	1.768067649	Up	2.17E-12	5.16E-12	Guanine nucleotide exchange factor OS=Culex quinquefasciatus GN=CpipJ_CPIJ017598 PE=4 SV=1
CCG026219.1	5121	303	3238	2.5424389	22.10406174	3.120026491	Up	1.37E-13	4.14E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007864 PE=4 SV=1
CCG025043.1	2664	625	3241	10.0811127	42.52994558	2.07682413	Up	0	0	AAEL005390-PA (Fragment) OS=Aedes aegypti GN=AAEL005390 PE=4 SV=1
CCG013979.1	939	181	3242	8.282771	120.6972666	3.865135692	Up	1.53E-13	4.55E-13	AAEL010422-PA OS=Aedes aegypti GN=AAEL010422 PE=4 SV=1
CCG002382.1	1758	130	3245	3.17751166	64.52764967	4.343948154	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012003 PE=4 SV=1
CCG023756.1	2688	1252	3247	20.014177	42.22824579	1.077186026	Up	5.40E-12	1.22E-11	AAEL006040-PA (Fragment) OS=Aedes aegypti GN=AAEL006040 PE=4 SV=1
CCG013169.3	2376	10367	3251	187.486212	47.83222129	-1.970729803	Down	0	0	AAEL000678-PA OS=Aedes aegypti GN=AAEL000678 PE=4 SV=1
CCG022945.1	1857	534	3258	12.3564019	61.33229172	2.311388161	Up	0	0	AAEL011217-PA OS=Aedes aegypti GN=AAEL011217 PE=4 SV=1
CCG026821.1	465	660	3261	60.9893007	245.1590119	2.007089715	Up	0	0	AAEL011104-PA OS=Aedes aegypti GN=AAEL011104 PE=4 SV=1

CCG025387.1	3471	1144	3261	14.1623095	32.84325571	1.213540593	Up	3.55E-12	8.26E-12	AGAP009115-PA (Fragment) OS=Anopheles gambiae GN=AGAP009115 PE=4 SV=4
CCG010316.1	4173	561	3263	5.77666454	27.33497654	2.242439515	Up	0	0	Sub1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ013695 PE=4 SV=1
CCG006483.1	3093	1163	3263	16.1570648	36.87968222	1.190661094	Up	4.90E-12	1.12E-11	AAEL0008800-PA OS=Aedes aegypti GN=AAEL008800 PE=4 SV=1
CCG024791.1	2242	519	3267	9.94705275	50.94053979	2.356473215	Up	0	0	AAEL001715-PA OS=Aedes aegypti GN=AAEL001715 PE=3 SV=1
CCG027692.1	1833	632	3268	14.8155332	62.32604937	2.072724724	Up	0	0	Adaptin, alpha/gamma/epsilon OS=Culex quinquefasciatus GN=CpipJ_CPIJ018379 PE=4 SV=1
CCG001293.3	4236	476	3276	4.82851598	27.03572113	2.485215082	Up	4.62E-13	1.21E-12	AAEL0006680-PA OS=Aedes aegypti GN=AAEL006680 PE=4 SV=1
CCG019023.1	2205	1089	3277	21.2217873	51.9538653	1.291684923	Up	3.37E-12	7.86E-12	AAEL0008854-PA OS=Aedes aegypti GN=AAEL008854 PE=4 SV=1
CCG011827.1	3057	27515	3278	386.755724	37.48551889	-3.367017365	Down	0	0	-
CCG000230.1	4064	1014	3280	10.7212871	28.21435724	1.395951366	Up	4.79E-12	1.09E-11	AAEL003368-PA OS=Aedes aegypti GN=AAEL003368 PE=4 SV=1
CCG015821.1	2498	488	3282	8.39440772	45.92996972	2.45193539	Up	9.43E-13	2.34E-12	AGAP005404-PA OS=Anopheles gambiae GN=AGAP005404 PE=4 SV=3
CCG012144.2	6568	1301	3282	8.51151412	17.46849336	1.037267481	Up	0	0	AAEL001296-PA OS=Aedes aegypti GN=AAEL001296 PE=4 SV=1
CCG008386.1	3636	622	3284	7.35070818	31.57397715	2.102780845	Up	0	0	AAEL014998-PA OS=Aedes aegypti GN=AAEL014998 PE=4 SV=1
CCG013869.1	2983	827	3284	11.9128295	38.4857462	1.691808096	Up	1.96E-12	4.67E-12	AAEL010483-PA OS=Aedes aegypti GN=AAEL010483 PE=4 SV=1
CCG017498.1	3878	1305	3285	14.459903	29.61267127	1.034156768	Up	8.11E-12	1.81E-11	AAEL0008020-PA OS=Aedes aegypti GN=AAEL008020 PE=4 SV=1
CCG026494.1	3198	204	3288	2.74103373	35.94209319	3.712882446	Up	6.59E-14	2.17E-13	Sub1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009759 PE=4 SV=1
CCG007000.1	2755	1083	3291	16.8915508	41.75959668	1.305805987	Up	4.78E-12	1.09E-11	AAEL003973-PA OS=Aedes aegypti GN=AAEL003973 PE=4 SV=1

CCG021885.1	3560	681	3292	8.21977227	32.32658627	1.975550836	Up	1.31E-14	4.67E-14	AAEL012236-PA OS=Aedes aegypti GN=AAEL012236 PE=4 SV=1
CCG004923.1	3141	325	3294	4.44608843	36.66111547	3.043642136	Up	1.19E-13	3.66E-13	AAEL011876-PA OS=Aedes aegypti GN=AAEL011876 PE=4 SV=1
CCG006466.1	9160	1130	3295	5.30085154	12.57505698	1.246268896	Up	2.84E-12	6.65E-12	-
CCG003027.1	1743	425	3309	10.4774166	66.36657363	2.663173749	Up	1.58E-12	3.80E-12	AAEL002978-PA OS=Aedes aegypti GN=AAEL002978 PE=4 SV=1
CCG005812.2	2070	678	3312	14.074145	55.93324284	1.990658698	Up	2.72E-13	7.58E-13	-
CCG016005.1	2114	911	3315	18.5172319	54.81867904	1.565799115	Up	0	0	AAEL001378-PA OS=Aedes aegypti GN=AAEL001378 PE=4 SV=1
CCG020491.1	34506	995	3316	1.23905657	3.359463449	1.43898878	Up	2.47E-12	5.84E-12	AAEL003548-PA (Fragment) OS=Aedes aegypti GN=AAEL003548 PE=4 SV=1
CCG003465.1	12532	462	3318	1.58410608	9.255630572	2.546662333	Up	0	0	AAEL008611-PA (Fragment) OS=Aedes aegypti GN=AAEL008611 PE=4 SV=1
CCG004055.1	2527	505	3320	8.58714522	45.92856307	2.419141152	Up	0	0	AAEL007637-PA OS=Aedes aegypti GN=AAEL007637 PE=3 SV=1
CCG005539.1	1827	177	3322	4.16291353	63.56398218	3.932544012	Up	0	0	AAEL008381-PA (Fragment) OS=Aedes aegypti GN=AAEL008381 PE=3 SV=1
CCG020156.1	3432	159	3327	1.99073071	33.88874907	4.089436399	Up	5.53E-14	1.85E-13	AAEL013690-PA OS=Aedes aegypti GN=AAEL013690 PE=4 SV=1
CCG017275.1	3942	762	3336	8.30617397	29.58417334	1.83256959	Up	5.84E-13	1.50E-12	AAEL010974-PA OS=Aedes aegypti GN=AAEL010974 PE=4 SV=1
CCG018837.1	2397	6912	3338	123.90772	48.6819891	-1.347806059	Down	0	0	AAEL010974-PA OS=Culex quinquefasciatus GN=CpipJ_CPIJ006896 PE=4 SV=1
CCG009439.1	13641	1106	3340	3.4839474	8.559537015	1.296809921	Up	2.74E-12	6.43E-12	AAEL004834-PA OS=Aedes aegypti GN=AAEL004834 PE=4 SV=1
CCG024407.1	513	13949	3341	1168.39148	227.6717401	-2.359496608	Down	0	0	Cytochrome c oxidase polypeptide VIIc OS=Aedes albopictus PE=4 SV=1
CCG011197.1	1891	354	3346	8.04404339	61.85636916	2.942929384	Up	0	0	AAEL013539-PA OS=Aedes aegypti GN=AAEL013539 PE=4 SV=1

CCG013063.1	1370	43159	3351	1353.6721	85.50743465	-3.984684642	Down	0	0	AAEL008166-PA OS=Aedes aegypti GN=AAEL008166 PE=3 SV=1
CCG006049.1	2403	182	3357	3.25447012	48.83684358	3.907475385	Up	0	0	AAEL011827-PA OS=Aedes aegypti GN=AAEL011827 PE=4 SV=1
CCG022910.1	3165	776	3357	10.5353915	37.07896844	1.815357183	Up	6.58E-13	1.67E-12	Impd1m-7 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002096 PE=4 SV=1
CCG019139.1	14829	1237	3360	3.58443332	7.92095286	1.143928937	Up	1.34E-13	4.07E-13	AAEL002306-PA OS=Aedes aegypti GN=AAEL002306 PE=4 SV=1
CCG018076.1	1118	10156	3363	390.340451	105.1562476	-1.892198412	Down	0	0	Triosephosphate isomerase OS=Aedes aegypti GN=AAEL002542 PE=3 SV=1
CCG014318.1	8199	5703	3368	29.8885713	14.36022395	-1.05751569	Down	6.68E-260	4.13E-259	AAEL006222-PA OS=Aedes aegypti GN=AAEL006222 PE=4 SV=1
CCG005843.1	387	37169	3369	4126.98208	304.326704	-3.761394337	Down	0	0	AAEL008844-PA OS=Aedes aegypti GN=AAEL008844 PE=4 SV=1
CCG000761.1	2871	1100	3370	16.4634999	41.03427124	1.317558271	Up	3.70E-12	8.58E-12	Atser1e-resistance protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002829 PE=4 SV=1
CCG028253.1	4014	217	3373	2.32297768	29.37575176	3.660578574	Up	1.47E-13	4.37E-13	AAEL006191-PA OS=Aedes aegypti GN=AAEL006191 PE=4 SV=1
CCG014196.1	3034	271	3373	3.83810088	38.86429385	3.339980765	Up	0	0	AAEL012057-PA OS=Aedes aegypti GN=AAEL012057 PE=4 SV=1
CCG005270.1	1160	941	3374	34.8573451	101.6803671	1.544506549	Up	0	0	AAEL012282-PA OS=Aedes aegypti GN=AAEL012282 PE=4 SV=1
CCG000490.1	3238	357	3375	4.73755258	36.43736384	2.943204727	Up	3.04E-13	8.40E-13	AAEL003152-PA OS=Aedes aegypti GN=AAEL003152 PE=4 SV=1
CCG024145.1	2226	857	3377	16.5431548	53.03418718	1.680688274	Up	1.00E-12	2.48E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009997 PE=4 SV=1
CCG011615.1	4901	610	3378	5.34820202	24.09489062	2.171601384	Up	3.62E-13	9.72E-13	DNA topoisomerase 2 (Fragment) OS=Aedes aegypti GN=AAEL012584 PE=3 SV=1
CCG004637.1	2724	815	3378	12.8562165	43.35134322	1.753610568	Up	1.33E-12	3.23E-12	AAEL007971-PA OS=Aedes aegypti GN=AAEL007971 PE=4 SV=1

CCG008891.1	2432	984	3381	17.3857808	48.59947935	1.483032999	Up	1.56E-12	3.75E-12	AAEL001187-PA OS=Aedes aegypti GN=AAEL001187 PE=4 SV=1
CCG020092.1	1224	1137	3391	39.9155133	96.84927821	1.278791734	Up	4.86E-12	1.11E-11	AAEL001379-PA OS=Aedes aegypti GN=AAEL001379 PE=4 SV=1
CCG015389.1	11921	8200	3397	29.5572371	9.96168662	-1.569049488	Down	0	0	AAEL0008773-PA OS=Aedes aegypti GN=AAEL008773 PE=4 SV=1
CCG018073.1	1491	116	3398	3.34305112	79.67017067	4.574802346	Up	1.16E-13	3.59E-13	AAEL002557-PA OS=Aedes aegypti GN=AAEL002557 PE=4 SV=1
CCG008652.1	6081	855	3402	6.04162524	19.55731912	1.69470002	Up	0	0	AAEL006022-PA (Fragment) OS=Aedes aegypti GN=AAEL006022 PE=4 SV=1
CCG028645.1	26274	7927	3402	12.9641884	4.526454197	-1.518078632	Down	0	0	AAEL002815-PA OS=Aedes aegypti GN=AAEL002815 PE=4 SV=1
CCG014798.2	5229	985	3405	8.09431796	22.76399549	1.491772372	Up	3.31E-12	7.72E-12	AAEL000343-PA OS=Aedes aegypti GN=AAEL000343 PE=4 SV=1
CCG000929.1	3356	226	3407	2.89367104	35.48952591	3.616420473	Up	2.04E-14	7.16E-14	AAEL013313-PA OS=Aedes aegypti GN=AAEL013313 PE=4 SV=1
CCG019773.1	2098	1384	3407	28.3460976	56.76970875	1.001971208	Up	3.46E-12	8.06E-12	AAEL005493-PA OS=Aedes aegypti GN=AAEL005493 PE=3 SV=1
CCG027373.1	761	0	3413	0.001	156.7839667	17.25841851	Up	4.44E-16	1.64E-15	-
CCG011390.1	917	1259	3419	58.9955244	130.3406197	1.143609344	Up	1.23E-12	2.99E-12	AAEL000670-PA OS=Aedes aegypti GN=AAEL000670 PE=3 SV=1
CCG017656.2	4877	731	3420	6.44061431	24.51451847	1.928366218	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002396 PE=4 SV=1
CCG025685.1	2085	558	3422	11.4998139	57.37516696	2.318815936	Up	2.34E-13	6.64E-13	AAEL011425-PA OS=Aedes aegypti GN=AAEL011425 PE=4 SV=1
CCG024366.2	1893	1314	3424	29.8268522	63.23145255	1.08403063	Up	1.36E-12	3.30E-12	AAEL007043-PA OS=Aedes aegypti GN=AAEL007043 PE=4 SV=1
CCG004908.1	743	18091	3425	1046.25231	161.1468344	-2.69878306	Down	0	0	-
CCG006392.1	1683	2	3430	0.05106326	71.24592355	10.44630606	Up	9.36E-05	0.00015362	AAEL000335-PA OS=Aedes aegypti GN=AAEL000335 PE=4 SV=1
CCG016652.1	2110	141	3431	2.87143724	56.84447754	4.307175262	Up	0	0	AAEL002843-PA OS=Aedes aegypti GN=AAEL002843 PE=4 SV=1

CCG020060.1	4234	1210	3433	12.2799667	28.34477188	1.206776014	Up	2.72E-12	6.41E-12	AAEL007987-PA OS=Aedes aegypti GN=AAEL007987 PE=4 SV=1
CCG012551.1	2895	678	3435	10.0633783	41.47899161	2.043266124	Up	2.72E-13	7.58E-13	AAEL002774-PA OS=Aedes aegypti GN=AAEL002774 PE=4 SV=1
CCG023536.1	2529	402	3436	6.83030182	47.49570541	2.797775834	Up	0	0	AAEL000118-PA OS=Aedes aegypti GN=AAEL000118 PE=4 SV=1
CCG017905.1	3090	1269	3436	17.6467939	38.8726987	1.139351171	Up	4.37E-12	1.00E-11	AAEL004808-PA OS=Aedes aegypti GN=AAEL004808 PE=4 SV=1
CCG028180.2	1160	1354	3438	50.1560523	103.6090996	1.04665501	Up	1.91E-12	4.57E-12	AAEL004278-PA OS=Aedes aegypti GN=AAEL004278 PE=4 SV=1
CCG021913.1	3216	495	3441	6.61381176	37.40405173	2.499640666	Up	1.42E-13	4.26E-13	AAEL03207 OS=Culex quinquefasciatus GN=CpipJ_CPIJ006841 PE=4 SV=1
CCG013306.1	171	1039	3448	261.085113	704.8896977	1.432877324	Up	2.45E-12	5.79E-12	40S ribosomal protein S29 OS=Aedes aegypti GN=AAEL007824 PE=4 SV=1
CCG028009.1	9597	964	3449	4.31622634	12.5634153	1.541386281	Up	1.62E-12	3.89E-12	histone-lysine N-methyltransferase ash1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001202 PE=4 SV=1
CCG025237.1	1769	451	3458	10.9549747	68.33562526	2.641051734	Up	1.51E-12	3.65E-12	AAEL003005-PA OS=Aedes aegypti GN=AAEL003005 PE=4 SV=1
CCG003927.2	3130	179	3468	2.4573746	38.73332391	3.97838561	Up	0	0	AAEL012224-PA OS=Aedes aegypti GN=AAEL012224 PE=4 SV=1
CCG006692.2	2345	943	3468	17.2795137	51.69948991	1.581087426	Up	1.51E-12	3.64E-12	AAEL005484-PA OS=Aedes aegypti GN=AAEL005484 PE=4 SV=1
CCG000728.1	582	881	3469	65.0452512	208.368148	1.67961912	Up	4.13E-13	1.09E-12	AAEL011090-PA OS=Aedes aegypti GN=AAEL011090 PE=2 SV=1
CCG014335.1	3339	1104	3471	14.2074235	36.34027514	1.354924397	Up	4.57E-12	1.05E-11	Pip1 interacting protein, pimt OS=Culex quinquefasciatus GN=CpipJ_CPIJ006162 PE=4 SV=1
CCG018337.1	3327	1104	3474	14.2586676	36.50287151	1.356170786	Up	4.57E-12	1.05E-11	Artemisinin-resistance protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002829 PE=4 SV=1

CCG002226.1	1299	777	3478	25.702451	93.59883496	1.864584633	Up	0	0	Discs overgrown protein kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ003503 PE=3 SV=1
CCG004101.1	3093	653	3487	9.0718515	39.41141646	2.119144671	Up	5.96E-13	1.53E-12	AAEL003108-PA OS=Aedes aegypti GN=AAEL003108 PE=4 SV=1
CCG024803.1	2972	709	3494	10.2508553	41.09832404	2.003335279	Up	2.32E-12	5.49E-12	AAEL013213-PA OS=Aedes aegypti GN=AAEL013213 PE=4 SV=1
CCG023657.1	3136	478	3496	6.54959603	38.97134426	2.572935865	Up	5.04E-13	1.31E-12	AAEL002290-PA OS=Aedes aegypti GN=AAEL002290 PE=4 SV=1
CCG018859.3	3744	690	3502	7.91910173	32.6986873	2.045824021	Up	8.73E-13	2.18E-12	AAEL011956-PA OS=Aedes aegypti GN=AAEL011956 PE=4 SV=1
CCG026987.1	2553	1372	3502	23.092235	47.95295153	1.054211806	Up	6.08E-12	1.37E-11	AAEL010359-PA OS=Aedes aegypti GN=AAEL010359 PE=3 SV=1
CCG016165.1	1590	1414	3502	38.2133363	76.99615425	1.010710167	Up	0	0	AAEL012026-PA OS=Aedes aegypti GN=AAEL012026 PE=2 SV=1
CCG028305.1	2797	12615	3507	193.801645	43.83220473	-2.144517667	Down	0	0	AAEL003606-PA OS=Aedes aegypti GN=AAEL003606 PE=4 SV=1
CCG003260.1	1068	18226	3512	733.301857	114.9564307	-2.673320019	Down	0	0	AAEL001490-PA OS=Aedes aegypti GN=AAEL001490 PE=3 SV=1
CCG021671.1	3108	141	3526	1.94939916	39.65987255	4.346578611	Up	0	0	AAEL013142-PA OS=Aedes aegypti GN=AAEL013142 PE=4 SV=1
CCG019517.1	2265	461	3530	8.7457164	54.48243576	2.639142732	Up	1.29E-13	3.95E-13	Mystery 43A OS=Culex quinquefasciatus GN=CpipJ_CPIJ001550 PE=4 SV=1
CCG003002.1	6105	622	3530	4.37791563	20.21338526	2.206994902	Up	0	0	AAEL007312-PA OS=Aedes aegypti GN=AAEL007312 PE=4 SV=1
CCG027862.1	1341	677	3531	21.6931472	92.04897486	2.085162286	Up	4.03E-13	1.07E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_00971 PE=4 SV=1
CCG005253.1	3782	1296	3533	14.7246896	32.65668742	1.149141235	Up	5.25E-12	1.19E-11	AAEL012750-PA OS=Aedes aegypti GN=AAEL012750 PE=4 SV=1
CCG002411.1	1669	1215	3535	31.2811429	74.04284505	1.243067105	Up	7.68E-13	1.93E-12	Signal transducer and activator of transcription OS=Aedes albopictus GN=STAT PE=2 SV=1

CCG005069.1	1512	577	3536	16.3978418	81.7542769	2.317788255	Up	4.04E-13	1.07E-12	AAEL011632-PA OS=Aedes aegypti GN=AAEL011632 PE=4 SV=1
CCG006450.1	1226	34449	3536	1207.39346	100.8258293	-3.581958721	Down	0	0	AAEL005170-PA OS=Aedes aegypti GN=AAEL005170 PE=4 SV=1
CCG001585.1	12067	1111	3541	3.95619252	10.25832917	1.374611231	Up	5.82E-12	1.32E-11	AAEL008134-PA OS=Aedes aegypti GN=AAEL008134 PE=3 SV=1
CCG010942.2	1773	1229	3546	29.7855634	69.91655354	1.231020824	Up	3.41E-13	9.25E-13	AAEL002983-PA (Fragment) OS=Aedes aegypti GN=AAEL002983 PE=4 SV=1
CCG025107.1	1422	18843	3546	569.39431	87.17443701	-2.70745103	Down	0	0	AAEL005610-PA OS=Aedes aegypti GN=AAEL005610 PE=4 SV=1
CCG000429.1	4512	434	3547	4.13317039	27.48160632	2.733145586	Up	0	0	AAEL005231-PA OS=Aedes aegypti GN=AAEL005231 PE=4 SV=1
CCG009126.1	616	835	3555	58.2463123	201.7478473	1.792314661	Up	0	0	AAEL009772-PA OS=Aedes aegypti GN=AAEL009772 PE=2 SV=1
CCG021405.1	2340	312	3559	5.72929795	53.16944745	3.214167202	Up	9.28E-14	2.93E-13	AAEL002771-PA (Fragment) OS=Aedes aegypti GN=AAEL002771 PE=4 SV=1
CCG016737.1	6147	1367	3563	9.55582027	20.2629478	1.084392444	Up	1.86E-12	4.45E-12	AAEL002392-PA OS=Aedes aegypti GN=AAEL002392 PE=4 SV=1
CCG013584.1	4875	1369	3566	12.0667829	25.57153128	1.08349746	Up	3.82E-12	8.84E-12	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC010545 PE=4 SV=1
CCG011382.1	1110	1028	3568	39.7953939	112.3703888	1.497588555	Up	1.50E-12	3.63E-12	AAEL003326-PA OS=Aedes aegypti GN=AAEL003326 PE=3 SV=1
CCG027364.1	2286	1020	3570	19.1728475	54.59363433	1.509668126	Up	1.22E-12	2.96E-12	AAEL005277-PA OS=Aedes aegypti GN=AAEL005277 PE=4 SV=1
CCG011237.1	2850	15495	3573	233.619662	43.82663962	-2.414281732	Down	0	0	AAEL012552-PA OS=Aedes aegypti GN=AAEL012552 PE=4 SV=1
CCG026280.1	925	294	3575	13.6574075	135.1090156	3.306368386	Up	0	0	60S acidic ribosomal protein P2 OS=Aedes aegypti GN=AAEL014583 PE=4 SV=1
CCG009221.1	2197	0	3576	0.001	56.90068172	15.79615832	Up	4.44E-16	1.64E-15	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1

CCG013784.1	2485	1075	3579	18.588517	50.34835918	1.437533088	Up	4.26E-12	9.83E-12	AAEL004167-PA OS=Aedes aegypti GN=AAEL004167 PE=4 SV=1
CCG027285.1	3475	1296	3580	16.0255471	36.01457003	1.168207073	Up	5.25E-12	1.19E-11	AAEL000981-PA OS=Aedes aegypti GN=AAEL000981 PE=4 SV=1
CCG009883.1	840	911	3582	46.6017003	149.0720802	1.677555582	Up	0	0	AAEL010954-PA OS=Aedes aegypti GN=AAEL010954 PE=4 SV=1
CCG019438.1	2748	1270	3587	19.8586474	45.63149155	1.200262452	Up	2.33E-12	5.50E-12	AAEL003785-PA OS=Aedes aegypti GN=AAEL003785 PE=4 SV=1
CCG000670.1	1143	672	3591	25.2630461	109.8295467	2.120165719	Up	0	0	AAEL009092-PB OS=Aedes aegypti GN=AAEL009092 PE=3 SV=1
CCG015783.1	3852	703	3596	7.84208812	32.63498527	2.05710963	Up	2.48E-13	6.98E-13	AAEL007690-PA OS=Aedes aegypti GN=AAEL007690 PE=4 SV=1
CCG002742.1	4000	447	3598	4.80186784	31.44496996	2.711161654	Up	3.44E-13	9.29E-13	AAEL001036-PA OS=Aedes aegypti GN=AAEL001036 PE=4 SV=1
CCG024776.1	836	36196	3599	1860.44559	150.4962178	-3.627849063	Down	0	0	Myosin light chain 2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017123 PE=4 SV=1
CCG013821.1	4038	445	3602	4.73539671	31.18368324	2.719234144	Up	0	0	AAEL011215-PA OS=Aedes aegypti GN=AAEL011215 PE=4 SV=1
CCG025131.2	15094	1428	3602	4.06524321	8.342368685	1.037115406	Up	1.05E-12	2.59E-12	AAEL014749-PB OS=Aedes aegypti GN=AAEL014749 PE=4 SV=1
CCG005477.1	2628	391	3603	6.39313784	47.92795708	2.906271343	Up	9.02E-13	2.25E-12	AAEL004493-PA OS=Aedes aegypti GN=AAEL004493 PE=4 SV=1
CCG010364.1	1851	8316	3603	193.050412	68.04682399	-1.504377903	Down	0	0	Actin filament light chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ010990 PE=4 SV=1
CCG010417.2	3062	1369	3608	19.2114849	41.19185584	1.100390096	Up	3.82E-12	8.84E-12	AAEL004936-PA OS=Aedes aegypti GN=AAEL004936 PE=4 SV=1
CCG018611.1	18286	871	3616	2.04673733	6.912891218	1.755963258	Up	0	0	AAEL004320-PA OS=Aedes aegypti GN=AAEL004320 PE=4 SV=1
CCG015651.1	1776	382	3620	9.24236409	71.25504612	2.946658358	Up	6.17E-13	1.58E-12	AGAP001945-PA OS=Anopheles gambiae GN=AgaP_AGAP001945 PE=4 SV=4
CCG000411.1	17281	641	3623	1.59386609	7.329080305	2.201101748	Up	0	0	AAEL007560-PA OS=Aedes aegypti GN=AAEL007560 PE=4 SV=1

CCG013682.1	1879	194	3634	4.43647074	67.60956774	3.929743066	Up	5.97E-13	1.53E-12	AAEL013775-PA OS=Aedes aegypti GN=AAEL013775 PE=4 SV=1
CCG005033.1	3045	398	3635	5.61640538	41.73180166	2.893428232	Up	7.64E-13	1.92E-12	AAEL011267-PA OS=Aedes aegypti GN=AAEL011267 PE=4 SV=1
CCG014352.1	5589	516	3636	3.96714673	22.74258263	2.519222433	Up	9.75E-14	3.07E-13	AAEL012442-PB (Fragment) OS=Aedes aegypti GN=AAEL012442 PE=4 SV=1
CCG021863.1	3045	25	3638	0.35278928	41.76624332	6.887386842	Up	9.32E-12	2.08E-11	Transcription factor btd OS=Culex quinquefasciatus GN=CpipJ_CPIJ018448 PE=4 SV=1
CCG006243.1	2989	303	3640	4.35591488	42.57214033	3.288861955	Up	1.37E-13	4.14E-13	AAEL003908-PA OS=Aedes aegypti GN=AAEL003908 PE=4 SV=1
CCG011818.1	7472	8617	3641	49.5543634	17.03467421	-1.540537739	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001569 PE=4 SV=1
CCG027327.1	8084	634	3646	3.36996682	15.76668445	2.22607302	Up	3.52E-13	9.51E-13	AAEL005215-PA OS=Aedes aegypti GN=AAEL005215 PE=4 SV=1
CCG003981.1	3177	410	3650	5.54535448	40.16295569	2.856513853	Up	1.25E-12	3.04E-12	AAEL011145-PA (Fragment) OS=Aedes aegypti GN=AAEL011145 PE=4 SV=1
CCG011935.1	6768	595	3654	3.77762886	18.87375049	2.320828264	Up	7.20E-13	1.81E-12	AAEL000079-PA OS=Aedes aegypti GN=AAEL000079 PE=4 SV=1
CCG027144.1	12297	650	3664	2.27131231	10.41612801	2.197221084	Up	0	0	AAEL008683-PA OS=Aedes aegypti GN=AAEL008683 PE=4 SV=1
CCG006404.1	4735	686	3665	6.22539344	27.05851835	2.119845921	Up	2.38E-13	6.73E-13	AAEL013609-PA (Fragment) OS=Aedes aegypti GN=AAEL013609 PE=4 SV=1
CCG019651.1	950	430	3666	19.4494588	134.9021502	2.794111425	Up	6.22E-13	1.59E-12	-
CCG028144.2	3933	706	3668	7.71335689	32.60283733	2.079566754	Up	0	0	AAEL000682-PA OS=Aedes aegypti GN=AAEL000682 PE=4 SV=1
CCG021692.1	3607	294	3685	3.5023848	35.71423618	3.350089763	Up	0	0	AAEL004286-PA OS=Aedes aegypti GN=AAEL004286 PE=4 SV=1
CCG015949.1	4592	1365	3687	12.7730156	28.06863381	1.135859669	Up	1.68E-12	4.04E-12	AAEL005706-PA OS=Aedes aegypti GN=AAEL005706 PE=4 SV=1

CCG019660.1	3417	525	3689	6.60202244	37.74102517	2.515153665	Up	2.05E-13	5.90E-13	AAEL002407-PA OS=Aedes aegypti GN=AAEL002407 PE=4 SV=1
CCG024832.1	1374	416	3696	13.0097595	94.03623796	2.853622527	Up	1.20E-14	4.28E-14	AAEL005890-PA OS=Aedes aegypti GN=AAEL005890 PE=4 SV=1
CCG011018.1	2043	252	3697	5.30023158	63.26027862	3.577172609	Up	5.42E-14	1.81E-13	AAEL012519-PA OS=Aedes aegypti GN=AAEL012519 PE=4 SV=1
CCG016301.1	2170	52	3698	1.02968949	59.57405876	5.854402995	Up	0	0	AAEL000584-PA OS=Aedes aegypti GN=AAEL000584 PE=4 SV=1
CCG026142.1	2136	253	3700	5.08957999	60.55506744	3.572629185	Up	3.13E-13	8.62E-13	AAEL001495-PA OS=Aedes aegypti GN=AAEL001495 PE=4 SV=1
CCG022091.1	5421	150	3702	1.18897993	23.8730014	4.327583693	Up	0	0	AAEL008709-PA OS=Aedes aegypti GN=AAEL008709 PE=4 SV=1
CCG014572.1	2613	647	3703	10.6396549	49.54094868	2.219170136	Up	0	0	AAEL011089-PA OS=Aedes aegypti GN=AAEL011089 PE=4 SV=1
CCG006028.1	2953	680	3712	9.89482544	43.94348912	2.150903263	Up	4.12E-13	1.09E-12	Heat shock 70 Cb OS=Aedes aegypti GN=Hsp70Cb PE=3 SV=1
CCG017851.1	1617	1293	3722	34.3598434	80.46673231	1.227668984	Up	5.13E-12	1.17E-11	AAEL008370-PA (Fragment) OS=Aedes aegypti GN=SCRB17 PE=4 SV=2
CCG010346.1	537	23	3723	1.84041694	242.3643658	7.041001149	Up	4.01E-11	8.68E-11	Galactose-specific C-type lectin OS=Culex quinquefasciatus GN=CpipJ_CPIJ012307 PE=4 SV=1
CCG000354.1	5258	405	3726	3.30976465	24.77263965	2.903947065	Up	2.34E-13	6.64E-13	AAEL007360-PA OS=Aedes aegypti GN=AAEL007360 PE=4 SV=1
CCG004592.1	14063	1222	3731	3.73384169	9.274644858	1.312631279	Up	6.16E-12	1.39E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_12310 PE=4 SV=1
CCG002166.1	3993	1469	3732	15.8082996	32.67325042	1.047427794	Up	3.08E-12	7.21E-12	AAEL005847-PA OS=Aedes aegypti GN=AAEL005847 PE=4 SV=1
CCG021696.1	2922	59	3737	0.86762982	44.7087886	5.687335003	Up	0	0	AAEL009395-PA OS=Aedes aegypti GN=AAEL009395 PE=4 SV=1
CCG011990.4	3003	1035	3738	14.8097487	43.51449836	1.554949005	Up	4.31E-12	9.93E-12	AAEL012778-PA OS=Aedes aegypti GN=AAEL012778 PE=4 SV=1
CCG015634.1	1158	37	3753	1.37295352	113.2974203	6.366688413	Up	5.17E-14	1.74E-13	AAEL011532-PA OS=Aedes aegypti GN=AAEL011532 PE=4 SV=1

CCG004950.1	2059	375	3758	7.82595944	63.80437305	3.02731577	Up	0	0	Suppressor of defective silencing OS=Culex quinquefasciatus GN=CpipJ_CPIJ016520 PE=4 SV_1
CCG023005.1	1981	252	3760	5.46611465	66.35190341	3.601550227	Up	5.42E-14	1.81E-13	AAEL011341-PA OS=Aedes aegypti GN=AAEL011341 PE=3 SV_1
CCG018563.1	2393	570	3765	10.2351645	55.0012169	2.425929244	Up	7.69E-13	1.93E-12	AAEL009612-PA OS=Aedes aegypti GN=AAEL009612 PE=4 SV_1
CCG024254.1	2971	660	3765	9.5456159	44.30087918	2.214425139	Up	0	0	AAEL008079-PA OS=Aedes aegypti GN=AAEL008079 PE=3 SV_1
CCG012703.2	5472	15517	3766	121.849666	24.05936958	-2.34043155	Down	0	0	Coagogen alpha-z (rv) chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ005295 PE=4 SV_1
CCG026893.1	528	331	3771	26.9374662	249.6736017	3.212357232	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005778 PE=4 SV_1
CCG010064.1	5128	771	3772	6.46054317	25.71423947	1.992840115	Up	0	0	AAEL006499-PA OS=Aedes aegypti GN=AAEL006499 PE=4 SV_1
CCG016613.1	3568	1467	3773	17.6672087	36.96681005	1.065156433	Up	6.83E-12	1.53E-11	AAEL006020-PA OS=Aedes aegypti GN=AAEL006020 PE=4 SV_1
CCG019742.1	1260	505	3777	17.2219968	104.7915963	2.605198695	Up	0	0	AAEL008600-PA OS=Aedes aegypti GN=AAEL008600 PE=3 SV_1
CCG009454.1	5033	572	3783	4.8835065	26.27601054	2.427756928	Up	0	0	AAEL006050-PA OS=Aedes aegypti GN=AAEL006050 PE=4 SV_1
CCG009687.2	3218	1502	3788	20.0561036	41.15038919	1.036864722	Up	5.54E-12	1.25E-11	AAEL001549-PA (Fragment) OS=Aedes aegypti GN=AAEL001549 PE=4 SV=1
CCG000952.2	4583	952	3797	8.92585367	28.96281407	1.698139724	Up	4.77E-13	1.24E-12	AAEL008171-PA OS=Aedes aegypti GN=AAEL008171 PE=4 SV_1
CCG021468.1	35208	1487	3800	1.81481468	3.773047368	1.055907975	Up	9.11E-12	2.04E-11	AAEL005822-PA OS=Aedes aegypti GN=AAEL005822 PE=3 SV_1
CCG022510.1	2820	906	3802	13.8051701	47.13169088	1.77148878	Up	0	0	AAEL013567-PA (Fragment) OS=Aedes aegypti GN=AAEL013567 PE=3 SV=1
CCG023266.1	6906	419	3808	2.60705456	19.27615377	2.886324533	Up	0	0	AAEL010978-PA OS=Aedes aegypti GN=AAEL010978 PE=4 SV_1

CCG024872.1	5408	845	3808	6.71402103	24.61559134	1.874323436	Up	0	0	AGAP001731-PA (Fragment) OS=Anopheles gambiae GN=AgaP_AGAP001731 PE=4 SV=5
CCG020887.1	1491	498	3811	14.3520643	89.35344921	2.638265165	Up	3.42E-13	9.27E-13	Goodpasture antigen-binding protein OS=Culex quinquefasciatus GN=Cq21.6P1.00098 PE=4 SV=1
CCG012802.1	11196	719	3814	2.75948903	11.90879489	2.109554372	Up	8.56E-13	2.14E-12	AAEL002788-PA OS=Aedes aegypti GN=AAEL002788 PE=4 SV=1
CCG007705.1	2257	933	3817	17.7628544	59.12084291	1.734803404	Up	7.48E-13	1.88E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_09750 PE=4 SV=1
CCG016859.1	2670	833	3819	13.4059135	50.00211947	1.899119723	Up	1.25E-12	3.05E-12	AAEL003646-PA OS=Aedes aegypti GN=AAEL003646 PE=4 SV=1
CCG018613.1	2052	25962	3821	543.655093	65.09530972	-3.06206616	Down	0	0	AAEL010602-PA OS=Aedes aegypti GN=AAEL010602 PE=4 SV=1
CCG005426.1	903	186	3826	8.85090879	148.1177928	4.064775551	Up	6.43E-13	1.64E-12	AAEL013628-PA OS=Aedes aegypti GN=AAEL013628 PE=3 SV=1
CCG013722.1	4749	119	3828	1.07673161	28.1786236	4.709870555	Up	0	0	DNA polymerase OS=Aedes aegypti GN=AAEL008247 PE=3 SV=1
CCG022472.1	3153	739	3829	10.0712445	42.45329583	2.075634595	Up	0	0	AAEL011111-PA (Fragment) OS=Aedes aegypti GN=AAEL011111 PE=4 SV=1
CCG016914.2	2388	834	3829	15.0070179	56.05328382	1.901161575	Up	4.58E-13	1.20E-12	AAEL011745-PA OS=Aedes aegypti GN=AAEL011745 PE=4 SV=1
CCG016666.1	13899	599	3835	1.85185057	9.645657344	2.380911873	Up	3.99E-13	1.06E-12	AAEL000416-PA OS=Aedes aegypti GN=AAEL000416 PE=4 SV=1
CCG019684.1	2105	874	3839	17.8411155	63.75526106	1.837338579	Up	0	0	AAEL012746-PA OS=Aedes aegypti GN=AAEL012746 PE=3 SV=1
CCG001777.1	884	600	3843	29.1649782	151.9735946	2.381511774	Up	0	0	-
CCG010584.1	1227	27	3846	0.94554428	109.5758211	6.856568749	Up	2.28E-12	5.41E-12	AAEL007584-PA OS=Aedes aegypti GN=AAEL007584 PE=4 SV=1
CCG024361.1	2200	895	3846	17.4808693	61.11342385	1.805712379	Up	9.98E-13	2.47E-12	-

CCG026924.1	2729	7596	3848	119.603556	49.29257934	-1.278817903	Down	0	0	AAEL005342-PA (Fragment) OS=Aedes aegypti GN=AAEL005342 PE=4 SV=1
CCG011408.1	3288	386	3851	5.04450047	40.94413742	3.020873574	Up	2.52E-13	7.09E-13	AAEL009636-PA OS=Aedes aegypti GN=AAEL009636 PE=4 SV=1
CCG007265.1	2128	6772	3851	136.743911	63.26331008	-1.112035646	Down	0	0	AAEL012419-PA OS=Aedes aegypti GN=AAEL012419 PE=4 SV=1
CCG017992.1	3075	966	3854	13.4987849	43.81437355	1.698574681	Up	4.38E-13	1.15E-12	AAEL003032-PA (Fragment) OS=Aedes aegypti GN=AAEL003032 PE=4 SV=1
CCG011783.1	3702	730	3855	8.47323238	36.40306779	2.103075695	Up	0	0	AAEL009269-PA OS=Aedes aegypti GN=AAEL009269 PE=4 SV=1
CCG006153.1	918	147	3859	6.8807745	146.9542375	4.416652188	Up	7.02E-14	2.29E-13	AAEL000005-PB OS=Aedes aegypti GN=AAEL000005 PE=4 SV=1
CCG008801.1	3717	148	3868	1.71092836	36.37842738	4.410231918	Up	0	0	AAEL014084-PA OS=Aedes aegypti GN=AAEL014084 PE=3 SV=1
CCG006660.1	1485	433	3872	12.5292223	91.15046981	2.862953226	Up	9.43E-13	2.34E-12	AGAP008118-PA OS=Anopheles gambiae GN=AGAP008118 PE=4 SV=4
CCG020338.1	3231	708	3874	9.41583785	41.91530926	2.154315892	Up	9.24E-13	2.30E-12	RNA polymerase II subunit A C- terminal domain phosphatase OS=Culex quinquefasciatus GN=CpipJ_CPIJ004583 PE=4 SV=1
CCG014360.1	2526	101	3878	1.71810895	53.66912008	4.965198809	Up	0	0	AAEL010153-PA OS=Aedes aegypti GN=AAEL010153 PE=4 SV=1
CCG026882.1	1863	1441	3879	33.2363862	72.78752313	1.130927644	Up	1.07E-12	2.62E-12	AAEL006021-PA OS=Aedes aegypti GN=AAEL006021 PE=4 SV=1
CCG023505.2	2567	935	3881	15.6512278	52.8527745	1.755703367	Up	1.31E-12	3.18E-12	AAEL005481-PA OS=Aedes aegypti GN=AAEL005481 PE=4 SV=1
CCG014983.1	1437	1260	3887	37.6770116	94.56007085	1.327546578	Up	1.31E-12	3.18E-12	AAEL006956-PA OS=Aedes aegypti GN=AAEL006956 PE=4 SV=1
CCG027111.1	23944	1450	3887	2.60215984	5.67502597	1.124917411	Up	1.87E-12	4.49E-12	AAEL008952-PA OS=Aedes aegypti GN=AAEL008952 PE=4 SV=1
CCG005143.1	663	1218	3888	78.9398744	205.0041932	1.37682729	Up	3.82E-12	8.85E-12	AAEL000951-PA OS=Aedes aegypti GN=AAEL000951 PE=3 SV=1

CCG004636.1	3809	1095	3889	12.3528116	35.69250154	1.530781569	Up	4.38E-12	1.01E-11	AAEL004289-PA OS=Aedes aegypti GN=AAEL004289 PE=4 SV=1
CCG020978.1	4407	598	3890	5.83070145	30.85720369	2.403865969	Up	0	0	Tyrosine-protein kinase receptor OS=Anopheles gambiae GN=INR PE=3 SV=3
CCG014228.2	2181	32039	3892	631.227569	62.38313306	-3.3389323	Down	0	0	AAEL003873-PA OS=Aedes aegypti GN=AAEL003873 PE=3 SV=1
CCG000030.1	4224	537	3894	5.46277166	32.2271614	2.560572095	Up	3.67E-13	9.84E-13	AAEL013724-PA (Fragment) OS=Aedes aegypti GN=AAEL013724 PE=4 SV=1
CCG015413.1	5634	554	3897	4.22528097	24.1804055	2.516719254	Up	9.41E-14	2.97E-13	AAEL013080-PA OS=Aedes aegypti GN=AAEL013080 PE=4 SV=1
CCG000730.2	3687	1424	3899	16.5958509	36.96835398	1.155468212	Up	4.81E-12	1.10E-11	AAEL011092-PA OS=Aedes aegypti GN=AAEL011092 PE=4 SV=1
CCG013691.1	3201	426	3900	5.71855887	42.59208979	2.896861992	Up	4.37E-13	1.15E-12	AAEL006559-PA OS=Aedes aegypti GN=AAEL006559 PE=4 SV=1
CCG020776.1	2697	980	3906	15.6137708	50.62922843	1.697151499	Up	1.36E-12	3.29E-12	Eukaryotic translation initiation factor 2C 2 OS=Culex quinquefasciatus GN=Cuq1.CPI006128 PE=4 SV=1
CCG009674.1	1509	712	3909	20.2746528	90.55792174	2.159163642	Up	0	0	AAEL002886-PA OS=Aedes aegypti GN=AAEL002886 PE=3 SV=1
CCG020520.1	2886	1346	3909	20.0406316	47.34993205	1.240434378	Up	4.85E-12	1.10E-11	AAEL008556-PA OS=Aedes aegypti GN=AAEL008556 PE=4 SV=1
CCG015800.2	3590	624	3920	7.46883409	38.17171168	2.353548924	Up	0	0	AAEL007484-PA OS=Aedes aegypti GN=AAEL007484 PE=4 SV=1
CCG014117.1	10584	843	3927	3.42247603	12.97063047	1.922136266	Up	2.68E-12	6.30E-12	AAEL009427-PA OS=Aedes aegypti GN=AAEL009427 PE=4 SV=1
CCG023025.2	5448	1181	3927	9.31484152	25.1984495	1.435731837	Up	1.60E-12	3.86E-12	AAEL006395-PA OS=Aedes aegypti GN=AAEL006395 PE=4 SV=1
CCG008348.1	6336	870	3928	5.9002003	21.67236603	1.877020827	Up	0	0	AAEL007242-PA (Fragment) OS=Aedes aegypti GN=AAEL007242 PE=4 SV=1
CCG016370.1	1810	15559	3935	369.373536	76.00045254	-2.281000594	Down	0	0	GL12416 OS=Drosophila persimilis GN=GL12416 PE=3 SV=1

CCG004620.1	3381	227	3938	2.88498366	40.71744866	3.819012112	Up	0	0	AAEL000154-PA OS=Aedes aegypti GN=AAEL000154 PE=4 SV=1
CCG010021.1	1803	269	3940	6.41090328	76.39246283	3.574830756	Up	0	0	AAEL012265-PA OS=Aedes aegypti GN=AAEL012265 PE=4 SV=1
CCG013047.1	4063	583	3943	6.16572859	33.92579013	2.460039126	Up	2.70E-13	7.54E-13	AAEL006062-PA (Fragment) OS=Aedes aegypti GN=AAEL006062 PE=4 SV=1
CCG015631.1	3558	1171	3944	14.1420908	38.75082732	1.45423168	Up	3.84E-12	8.89E-12	AAEL007847-PA (Fragment) OS=Aedes aegypti GN=AAEL007847 PE=4 SV=1
CCG017863.3	5949	1200	3954	8.66762171	23.23500191	1.422591669	Up	4.83E-13	1.26E-12	AAEL001296-PA OS=Aedes aegypti GN=AAEL001296 PE=4 SV=1
CCG024595.1	4383	495	3956	4.85284477	31.55257653	2.7008552	Up	1.42E-13	4.26E-13	AAEL005532-PA (Fragment) OS=Aedes aegypti GN=AAEL005532 PE=4 SV=1
CCG002067.1	5451	874	3956	6.88966209	25.37056373	1.880650445	Up	0	0	AAEL008814-PA OS=Aedes aegypti GN=AAEL008814 PE=4 SV=1
CCG022488.2	1203	183	3957	6.53654317	114.987449	4.136804716	Up	3.05E-13	8.41E-13	AAEL012183-PA OS=Aedes aegypti GN=AAEL012183 PE=4 SV=1
CCG018981.1	2516	1061	3958	18.1203849	54.9939823	1.60166016	Up	3.27E-12	7.64E-12	AAEL010584-PA OS=Aedes aegypti GN=AAEL010584 PE=4 SV=1
CCG010930.1	4906	1088	3958	9.52936634	28.2031919	1.56540626	Up	2.82E-12	6.61E-12	AAEL008244-PA (Fragment) OS=Aedes aegypti GN=AAEL008244 PE=4 SV=1
CCG008896.1	2927	1282	3958	18.8203621	47.27190279	1.328688554	Up	3.23E-12	7.55E-12	AAEL000802-PA (Fragment) OS=Aedes aegypti GN=AAEL000802 PE=4 SV=1
CCG024011.1	2166	761	3960	15.0969382	63.91263897	2.081845275	Up	0	0	AAEL008938-PA OS=Aedes aegypti GN=AAEL008938 PE=4 SV=1
CCG016977.1	2025	1072	3960	22.7474348	68.36285235	1.587508728	Up	4.51E-12	1.04E-11	AAEL006589-PA OS=Aedes aegypti GN=AAEL006589 PE=4 SV=1
CCG001159.1	3932	1199	3960	13.1029277	35.20721669	1.425981975	Up	4.07E-12	9.41E-12	AAEL008210-PA OS=Aedes aegypti GN=AAEL008210 PE=4 SV=1

CCG011598.1	6139	941	3961	6.58649947	22.55574756	1.775911277	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004311 PE=4 SV=1
CCG009754.1	4368	163	3965	1.60349513	31.73295957	4.3066902	Up	1.80E-13	5.24E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013634 PE=4 SV=1
CCG015338.1	8340	1333	3966	6.86794439	16.62404385	1.275321101	Up	4.28E-12	9.87E-12	AAEL002037-PA OS=Aedes aegypti GN=AAEL002037 PE=4 SV=1
CCG006538.1	2922	1530	3968	22.499553	47.47243061	1.077193577	Up	5.34E-12	1.21E-11	AAEL007817-PA OS=Aedes aegypti GN=AAEL007817 PE=4 SV=1
CCG013477.1	2196	318	3972	6.22239326	63.23054433	3.345080156	Up	1.20E-13	3.69E-13	AAEL014701-PA OS=Aedes aegypti GN=AAEL014701 PE=4 SV=1
CCG016115.2	2829	737	3987	11.194307	49.2678153	2.137880285	Up	0	0	AAEL000509-PA OS=Aedes aegypti GN=AAEL000509 PE=4 SV=1
CCG004495.1	1716	1076	3988	26.9437264	81.24336117	1.592300536	Up	5.36E-12	1.21E-11	AAEL007787-PA OS=Aedes aegypti GN=AAEL007787 PE=4 SV=1
CCG007420.1	9238	317	3993	1.47449728	15.11024022	3.35723153	Up	2.22E-13	6.35E-13	2-Cys thioredoxin peroxidase OS=Aedes aegypti GN=TPx PE=2 SV=1
CCG014824.1	1468	333	3994	9.74722182	95.1112789	3.286553454	Up	4.41E-13	1.16E-12	AAEL010248-PA (Fragment) OS=Aedes aegypti GN=AAEL010248 PE=3 SV=1
CCG025015.2	4651	8188	3996	75.6474279	30.03510513	-1.332641315	Down	0	0	AAEL007651-PA OS=Aedes aegypti GN=AAEL007651 PE=4 SV=1
CCG023516.4	27883	1126	3999	1.73524804	5.013741305	1.530745658	Up	5.84E-12	1.32E-11	AAEL007417-PA OS=Aedes aegypti GN=AAEL007417 PE=4 SV=1
CCG026840.1	1523	447	4001	12.611603	91.83720641	2.864327096	Up	3.44E-13	9.29E-13	AAEL003664-PA OS=Aedes aegypti GN=AAEL003664 PE=4 SV=1
CCG003944.1	1095	6865	4006	269.394729	127.8930199	-1.074784095	Down	0	0	AAEL015665-PA OS=Aedes aegypti GN=AAEL015665 PE=4 SV=1
CCG021140.1	902	1604	4008	76.4118119	155.3356689	1.023521571	Up	6.44E-12	1.45E-11	AAEL004450-PA OS=Aedes aegypti GN=AAEL004450 PE=3 SV=1
CCG024735.1	1770	7	4012	0.1699368	79.23876068	8.865064172	Up	4.31E-06	7.70E-06	AAEL003404-PA OS=Aedes aegypti GN=AAEL003404 PE=4 SV=1

CCG011646.1	1860	489	4012	11.2968818	75.4046271	2.738728439	Up	1.91E-13	5.52E-13	AAEL004832-PA OS=Aedes aegypti GN=AAEL004832 PE=4 SV=1
CCG024915.1	1650	6759	4025	176.019658	85.27700849	-1.045507827	Down	6.22E-302	3.93E-301	AAEL006776-PA OS=Aedes aegypti GN=AAEL006776 PE=4 SV=1
CCG023458.1	20124	24793	4025	52.9392084	6.992002783	-2.920558959	Down	0	0	AAEL002565-PA OS=Aedes aegypti GN=AAEL002565 PE=4 SV=1
CCG010440.1	6617	622	4035	4.03916804	21.31731098	2.399895392	Up	0	0	AAEL007586-PA OS=Aedes aegypti GN=AAEL007586 PE=4 SV=1
CCG022225.1	804	831	4035	44.4127481	175.4435905	1.981961495	Up	5.11E-15	1.84E-14	AAEL009244-PA OS=Aedes aegypti GN=AAEL009244 PE=3 SV=1
CCG015078.1	2202	384	4057	7.49335971	64.40768795	3.10354829	Up	1.03E-12	2.55E-12	AAEL012441-PA OS=Aedes aegypti GN=AAEL012441 PE=4 SV=1
CCG000789.1	3171	418	4060	5.66425388	44.75894156	2.982218084	Up	6.74E-13	1.70E-12	AAEL002903-PA OS=Aedes aegypti GN=AAEL002903 PE=4 SV=1
CCG016815.1	7001	939	4062	5.76325965	20.28289105	1.81530638	Up	2.55E-13	7.16E-13	AAEL008662-PA (Fragment) OS=Aedes aegypti GN=AAEL008662 PE=4 SV=1
CCG011133.1	6672	252	4069	1.6229576	21.31972845	3.715491845	Up	5.42E-14	1.81E-13	AAEL014706-PA OS=Aedes aegypti GN=AAEL014706 PE=4 SV=1
CCG003706.1	3034	891	4069	12.6189959	46.88372715	1.893490147	Up	0	0	AAEL005476-PA OS=Aedes aegypti GN=AAEL005476 PE=4 SV=1
CCG013723.1	1308	1124	4075	36.9250625	108.9105335	1.560471228	Up	3.38E-12	7.88E-12	AAEL008257-PA OS=Aedes aegypti GN=AAEL008257 PE=2 SV=1
CCG009267.1	1620	1116	4082	29.6013727	88.08622579	1.573252359	Up	3.58E-12	8.33E-12	Transiocon-associated protein subunit alpha OS=Culex quinquefasciatus GN=CpipJ_CPIJ018234 PE=4 SV=1
CCG021799.1	5775	914	4083	6.80075107	24.71595568	1.861676701	Up	0	0	AAEL011187-PA OS=Aedes aegypti GN=AAEL011187 PE=4 SV=1
CCG010962.1	2467	1366	4085	23.792727	57.88591837	1.282691799	Up	2.64E-12	6.21E-12	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC003691 PE=4 SV=1

CCG014372.1	1830	461	4087	10.8246162	78.07348479	2.850516791	Up	1.29E-13	3.95E-13	Alkaline phosphatase OS=Aedes aegypti GN=AAEL009077 PE=3 SV=1
CCG021143.1	2996	670	4092	9.60938658	47.74675185	2.312886348	Up	8.60E-13	2.15E-12	AAEL004477-PA OS=Aedes aegypti GN=AAEL004477 PE=3 SV=1
CCG001107.1	2618	1047	4094	17.1846112	54.66737399	1.669562864	Up	5.84E-12	1.32E-11	AAEL009307-PA (Fragment) OS=Aedes aegypti GN=AAEL009307 PE=4 SV=1
CCG003786.1	14045	1473	4097	4.50654461	10.19751228	1.178123666	Up	7.16E-12	1.60E-11	GTP-binding protein SARZ OS=Culex quinquefasciatus GN=CpipJ_CPIJ012024 PE=3 SV=1
CCG020553.1	1131	63552	4099	2414.51156	126.6967078	-4.252280429	Down	0	0	AAEL005961-PA OS=Aedes aegypti GN=AAEL005961 PE=3 SV=1
CCG014018.1	1373	27074	4104	847.314344	104.4929118	-3.019492212	Down	0	0	AAEL004439-PA OS=Aedes aegypti GN=AAEL004439 PE=4 SV=1
CCG007520.1	22321	1578	4107	3.03777793	6.432222692	1.082300946	Up	4.46E-13	1.17E-12	AAEL008507-PA OS=Aedes aegypti GN=AAEL008507 PE=3 SV=1
CCG014049.1	8265	1579	4107	8.20922093	17.37128163	1.08138698	Up	4.25E-12	9.80E-12	AAEL011479-PA OS=Aedes aegypti GN=AAEL011479 PE=4 SV=1
CCG023374.1	2601	1341	4115	22.1539462	55.30692384	1.319896397	Up	2.45E-12	5.78E-12	AAEL010599-PA OS=Aedes aegypti GN=AAEL010599 PE=4 SV=1
CCG001726.1	3026	615	4116	8.73310865	47.5506501	2.44489787	Up	0	0	AAEL003293-PA OS=Aedes aegypti GN=AAEL003293 PE=4 SV=1
CCG001673.1	5895	1518	4119	11.06498	24.42631756	1.142435539	Up	2.23E-12	5.30E-12	AAEL006687-PA (Fragment) OS=Aedes aegypti GN=AAEL006687 PE=4 SV=1
CCG023250.1	1746	827	4122	20.3527895	82.53036475	2.019698474	Up	1.96E-12	4.67E-12	AAEL010841-PA OS=Aedes aegypti GN=AAEL010841 PE=4 SV=1
CCG003509.1	2930	1128	4125	16.5426146	49.21600399	1.572940255	Up	5.47E-12	1.24E-11	AAEL012286-PA OS=Aedes aegypti GN=AAEL012286 PE=4 SV=1
CCG019710.1	6042	228	4128	1.62149942	23.88410568	3.880650445	Up	1.05E-13	3.28E-13	Nuclear pore complex protein Nup205 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003755 PE=4 SV=1

CCG015609.1	8919	1288	4128	6.20529411	16.17981461	1.382623581	Up	7.16E-12	1.61E-11	Nuclear receptor co-repressor 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002507 PE=4 SV=1
CCG005820.1	1158	706	4129	26.1974375	124.6482943	2.250365534	Up	0	0	AAEL000970-PA OS=Aedes aegypti GN=AAEL000970 PE=4 SV=1
CCG021289.1	1851	1036	4129	24.0500514	77.98094262	1.697081619	Up	3.37E-12	7.87E-12	AAEL011063-PA OS=Aedes aegypti GN=AAEL011063 PE=4 SV=1
CCG007743.1	2064	1067	4129	22.2135208	69.93349069	1.654545446	Up	2.93E-12	6.87E-12	AAEL006057-PA OS=Aedes aegypti GN=AAEL006057 PE=4 SV=1
CCG026099.1	13519	1306	4129	4.15108169	10.67702676	1.362950725	Up	6.89E-13	1.74E-12	AAEL008105-PA OS=Aedes aegypti GN=AAEL008105 PE=4 SV=1
CCG009285.1	2983	1516	4136	21.8377867	48.47047695	1.150279636	Up	2.16E-12	5.15E-12	AAEL013845-PA OS=Aedes aegypti GN=AAEL013845 PE=4 SV=1
CCG001405.1	4113	483	4138	5.04604469	35.17076326	2.801151755	Up	0	0	AAEL003008-PA OS=Aedes aegypti GN=AAEL003008 PE=4 SV=1
CCG007055.1	5343	1218	4139	9.79545887	27.08072386	1.467081319	Up	3.82E-12	8.85E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000283 PE=4 SV=1
CCG023329.2	14223	1532	4139	4.62839298	10.17312153	1.136179155	Up	1.74E-12	4.18E-12	AAEL005174-PA OS=Aedes aegypti GN=AAEL005174 PE=4 SV=1
CCG002937.1	4964	1513	4140	13.0969397	29.15537184	1.154531984	Up	0	0	AAEL003736-PA OS=Aedes aegypti GN=AAEL003736 PE=4 SV=1
CCG024477.1	966	12	4145	0.53378552	150.0021296	8.134507089	Up	1.30E-07	2.49E-07	Long form D7 salivary protein D7I2 OS=Aedes albopictus PE=2 SV=1
CCG027348.1	3087	730	4145	10.1612913	46.93944192	2.207716936	Up	0	0	AAEL003968-PA OS=Aedes aegypti GN=AAEL003968 PE=4 SV=1
CCG012743.1	1028	82	4170	3.42754692	141.8054612	5.370592868	Up	0	0	AAEL010094-PA OS=Aedes aegypti GN=AAEL010094 PE=3 SV=1
CCG016741.1	7746	1645	4172	9.12538258	18.82854773	1.044964778	Up	3.83E-12	8.87E-12	AAEL008642-PA OS=Aedes aegypti GN=AAEL008642 PE=4 SV=1
CCG027742.1	513	89	4187	7.45478827	285.3222317	5.258280976	Up	0	0	AAEL009844-PA OS=Aedes aegypti GN=AAEL009844 PE=4 SV=1
CCG005192.1	2253	802	4187	15.2959286	64.96684636	2.08655598	Up	7.06E-13	1.78E-12	AAEL000706-PA OS=Aedes aegypti GN=AAEL000706 PE=3 SV=1

CCG021754.2	4639	858	4190	7.94740942	31.57473155	1.990213895	Up	1.49E-14	5.27E-14	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007358 PE=4 SV=1
CCG009756.1	3002	291	4191	4.1652874	48.80417653	3.550516667	Up	6.00E-14	1.99E-13	AAEL000196-PA OS=Aedes aegypti GN=AAEL006196 PE=4 SV=1
CCG019170.1	2595	625	4194	10.3491654	56.49904153	2.44871197	Up	0	0	AAEL015099-PA OS=Aedes aegypti GN=AAEL015099 PE=4 SV=1
CCG000352.1	1785	415	4195	9.99016239	82.15684653	3.039800773	Up	0	0	AAEL004613-PA OS=Aedes aegypti GN=AAEL004613 PE=4 SV=1
CCG024232.1	2840	293	4197	4.43314516	51.66193226	3.542699097	Up	4.19E-13	1.11E-12	Ribonucleoside-diphosphate reductase OS=Aedes aegypti GN=RNR1 PE=2 SV=1
CCG028275.1	2037	952	4198	20.0820753	72.04459788	1.842981892	Up	4.77E-13	1.24E-12	AAEL006372-PA OS=Aedes aegypti GN=AAEL006372 PE=4 SV=1
CCG009142.1	737	32748	4201	1909.32547	199.2669209	-3.260288906	Down	0	0	ATP synthase-coupling factor 6, mitochondrial OS=Aedes albopictus PE=3 SV=1
CCG004691.1	4947	812	4203	7.0530472	29.70075546	2.074181028	Up	1.91E-12	4.56E-12	AAEL008306-PA OS=Aedes aegypti GN=AAEL008306 PE=4 SV=1
CCG020752.1	903	1365	4209	64.95425	162.9450575	1.326889762	Up	1.68E-12	4.04E-12	AAEL004129-PA OS=Aedes aegypti GN=AAEL004129 PE=4 SV=1
CCG012468.1	2085	1429	4211	29.4502402	70.60398249	1.261470163	Up	2.28E-12	5.41E-12	AAEL012821-PA OS=Aedes aegypti GN=AAEL012821 PE=4 SV=1
CCG010793.1	2315	454	4212	8.42689396	63.60443273	2.916054438	Up	3.16E-13	8.66E-13	AAEL006752-PA OS=Aedes aegypti GN=AAEL006752 PE=4 SV=1
CCG004698.1	38283	12182	4217	13.6733617	3.850770659	-1.828148879	Down	0	0	AAEL001504-PA OS=Aedes aegypti GN=AAEL001504 PE=4 SV=1
CCG011203.1	1095	280	4220	10.9876947	134.7250484	3.61605747	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010850 PE=4 SV=1
CCG002241.1	4737	140	4224	1.26995205	31.17242159	4.617424306	Up	0	0	Microtubule binding protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012473 PE=4 SV=1

CCG019558.1	2652	1106	4227	17.9202589	55.7196968	1.636595931	Up	2.74E-12	6.43E-12	Phosphatidylinositol 3-kinase regulatory subunit OS=Culex quinquefasciatus GN=CpipJ_CPIJ011236 PE=4 SV=1
CCG025858.1	7266	1137	4232	6.72400059	20.36105523	1.598420577	Up	4.86E-12	1.11E-11	AAEL000693-PA OS=Aedes aegypti GN=AAEL000693 PE=4 SV=1
CCG009649.3	2304	610	4235	11.3765356	64.25707558	2.497794026	Up	3.62E-13	9.71E-13	M-kinase phosphoprotein 11 OS=Culex quinquefasciatus GN=CpipJ_CPIJ011412 PE=4 SV=1
CCG026982.1	5388	612	4241	4.88074937	27.5163422	2.495114129	Up	6.59E-13	1.68E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_16871 PE=4 SV=1
CCG020637.1	1917	991	4246	22.2133579	77.42975648	1.801460613	Up	5.56E-12	1.26E-11	AAEL001432-PA OS=Aedes aegypti GN=AAEL001432 PE=3 SV=1
CCG023730.1	2289	931	4254	17.4769869	64.96833088	1.894280164	Up	9.75E-13	2.41E-12	AAEL013644-PA OS=Aedes aegypti GN=AAEL013644 PE=4 SV=1
CCG021377.1	3294	657	4256	8.57046619	45.16770672	2.397846079	Up	0	0	AAEL008741-PA OS=Aedes aegypti GN=AAEL008741 PE=4 SV=1
CCG012962.1	11589	11258	4262	41.7424517	12.85634443	-1.699034851	Down	0	0	AAEL000405-PA (Fragment) OS=Aedes aegypti GN=AAEL000405 PE=4 SV=1
CCG009805.2	3630	375	4263	4.43902217	41.05430685	3.209219758	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_19518 PE=4 SV=1
CCG016683.1	1762	1233	4264	30.0690595	84.59823618	1.492347842	Up	3.53E-12	8.21E-12	Adenylate kinase OS=Glossina morsitans morsitans PE=2 SV=1
CCG008919.1	2336	931	4266	17.1253523	63.84075715	1.898344097	Up	9.75E-13	2.41E-12	AISZC17 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008428 PE=4 SV=1
CCG025989.2	633	31910	4274	2166.13623	236.0374011	-3.198036592	Down	0	0	AAEL010823-PA OS=Aedes aegypti GN=AAEL010823 PE=3 SV=1
CCG019046.2	4554	1739	4282	16.4085131	32.87029889	1.002340067	Up	4.99E-12	1.13E-11	AAEL009142-PA OS=Aedes aegypti GN=AAEL009142 PE=4 SV=1
CCG019454.1	8531	1544	4290	7.77696287	17.57953433	1.176618099	Up	4.92E-12	1.12E-11	AAEL011640-PA OS=Aedes aegypti GN=AAEL011640 PE=4 SV=1

CCG008892.2	4737	809	4291	7.33850861	31.66687052	2.109415497	Up	1.25E-12	3.04E-12	AAEL001165-PA OS=Aedes aegypti GN=AAEL001165 PE=4 SV=1
CCG027599.1	3597	496	4292	5.92521222	41.71279508	2.815551254	Up	6.79E-13	1.71E-12	AAEL010913-PA (Fragment) OS=Aedes aegypti GN=AAEL010913 PE=4 SV=1
CCG016653.1	2329	1149	4292	21.1988944	64.42289562	1.603584482	Up	3.86E-12	8.94E-12	AAEL002834-PA OS=Aedes aegypti GN=AAEL002834 PE=4 SV=1
CCG020203.1	28922	9136	4292	13.573456	5.187778297	-1.387599371	Down	0	0	AAEL013633-PA OS=Aedes aegypti GN=AAEL013633 PE=3 SV=1
CCG007967.1	6761	1286	4296	8.17321087	22.2128024	1.442416554	Up	6.57E-12	1.48E-11	AAEL002235-PA OS=Aedes aegypti GN=AAEL002235 PE=4 SV=1
CCG022179.1	4626	846	4298	7.85827831	32.4796095	2.047249118	Up	0	0	AAEL007447-PA OS=Aedes aegypti GN=AAEL007447 PE=4 SV=2
CCG016423.1	1011	508	4305	21.5911228	148.8579441	2.785426039	Up	6.20E-13	1.58E-12	AAEL007046-PA OS=Aedes aegypti GN=AAEL007046 PE=3 SV=1
CCG001369.2	5760	530	4308	3.95381239	26.14587784	2.725267189	Up	0	0	AAEL004290-PA (Fragment) OS=Aedes aegypti GN=AAEL004290 PE=3 SV=1
CCG027631.1	1306	31420	4309	1033.77417	115.3408994	-3.163944973	Down	0	0	AAEL010143-PA OS=Aedes aegypti GN=AAEL010143 PE=3 SV=1
CCG025106.2	2059	22802	4316	475.860072	73.27825282	-2.699080397	Down	0	0	AAEL005617-PA OS=Aedes aegypti GN=UGP PE=2 SV=1
CCG007833.1	5376	178	4321	1.42273303	28.09797506	4.303729288	Up	1.15E-13	3.57E-13	AAEL008709-PA OS=Aedes aegypti GN=AAEL008709 PE=4 SV=1
CCG024098.1	2371	32	4326	0.57993737	63.78300519	6.781131154	Up	1.77E-13	5.17E-13	AAEL013813-PA OS=Aedes aegypti GN=AAEL013813 PE=4 SV=1
CCG020586.1	2190	1024	4334	20.0917846	69.18227011	1.783796642	Up	2.58E-12	6.08E-12	AAEL007397-PC OS=Aedes aegypti GN=e75B PE=2 SV=1
CCG006899.1	4362	389	4336	3.8320098	34.74990557	3.1808359	Up	0	0	AAEL006963-PA OS=Aedes aegypti GN=AAEL006963 PE=4 SV=1
CCG011655.1	2706	99	4338	1.57206346	56.04176077	5.155770921	Up	0	0	AAEL004831-PA OS=Aedes aegypti GN=AAEL004831 PE=4 SV=1
CCG011548.1	1998	326	4339	7.01107782	75.91789936	3.43673192	Up	6.17E-13	1.58E-12	AAEL002981-PA OS=Aedes aegypti GN=AAEL002981 PE=4 SV=1

CCG011658.1	5190	489	4342	4.04859349	29.24640419	2.852766559	Up	1.91E-13	5.52E-13	AAEL000386-PA OS=Aedes aegypti GN=AAEL000386 PE=4 SV=1
CCG023117.1	5154	716	4345	5.96940822	29.47103465	2.303637888	Up	0	0	AAEL014260-PA OS=Aedes aegypti GN=AAEL014260 PE=4 SV=1
CCG019844.1	7719	519	4346	2.88914267	19.68242918	2.768194935	Up	0	0	AAEL012147-PA OS=Aedes aegypti GN=AAEL012147 PE=4 SV=1
CCG005931.1	3515	954	4353	11.6623405	43.29256865	1.892262053	Up	2.67E-12	6.29E-12	AAEL007201-PA (Fragment) OS=Aedes aegypti GN=AAEL007201 PE=4 SV=1
CCG018646.1	3432	1723	4358	21.5725095	44.39049247	1.041056699	Up	0	0	AAEL013920-PA OS=Aedes aegypti GN=AAEL013920 PE=4 SV=1
CCG024363.1	2181	1257	4366	24.7652253	69.98066776	1.498640684	Up	1.92E-12	4.59E-12	AAEL007042-PA OS=Aedes aegypti GN=AAEL007042 PE=4 SV=1
CCG007013.1	3336	953	4384	12.2752269	45.94037331	1.904012883	Up	1.54E-12	3.70E-12	AAEL003901-PA OS=Aedes aegypti GN=AAEL003901 PE=4 SV=1
CCG000010.1	4673	1593	4386	14.6481462	32.81125656	1.163472749	Up	2.15E-12	5.13E-12	AAEL001273-PA OS=Aedes aegypti GN=AAEL001273 PE=4 SV=1
CCG000489.1	4162	763	4390	7.87744054	36.87333855	2.226779181	Up	4.81E-13	1.25E-12	DNA-directed RNA polymerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007655 PE=3 SV=1
CCG028292.1	4896	917	4397	8.04804874	31.39533149	1.963839097	Up	0	0	AAEL009893-PA OS=Aedes aegypti GN=AAEL009893 PE=4 SV=1
CCG027126.1	40417	1193	4397	1.26834979	3.803140831	1.584238693	Up	2.22E-12	5.28E-12	AGAP013535-PA OS=Anopheles gambiae GN=AgaP_AGAP013535 PE=4 SV=1
CCG022444.1	9367	770	4400	3.53226173	16.42109723	2.216886377	Up	0	0	AAEL010502-PA OS=Aedes aegypti GN=AAEL010502 PE=4 SV=1
CCG012397.1	3091	1486	4403	20.6577242	49.79660066	1.269365932	Up	5.90E-12	1.33E-11	Uridine kinase OS=Aedes aegypti GN=AAEL011273 PE=3 SV=1
CCG026861.1	2289	558	4407	10.4749288	67.30499158	2.683773073	Up	2.34E-13	6.64E-13	AAEL001129-PA OS=Aedes aegypti GN=AAEL001129 PE=4 SV=1
CCG027478.1	1177	1722	4418	62.8665106	131.2197679	1.06162148	Up	1.20E-12	2.92E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_18279 PE=4 SV=1

CCG013376.1	801	313	4432	16.79092	193.4270695	3.526036523	Up	0	0	AAEL005312-PE OS=Aedes aegypti PE=2 SV=1
CCG019932.1	6780	1480	4433	9.37982407	22.85693819	1.28499939	Up	6.02E-12	1.36E-11	Ubiquitin carboxyl-terminal hydrolase (Fragment) OS=Aedes aegypti GN=AAEL009824 PE=3 SV=1
CCG002687.2	2382	1140	4438	20.5648604	65.13217142	1.663189047	Up	4.59E-12	1.05E-11	AAEL001919-FA OS=Aedes aegypti GN=AAEL001919 PE=4 SV=1
CCG004630.1	820	476	4442	24.9434069	189.3715432	2.924489118	Up	4.62E-13	1.21E-12	AAEL008441-FA OS=Aedes aegypti GN=AAEL008441 PE=4 SV=1
CCG024905.1	3222	800	4447	10.6690837	48.24936586	2.177073703	Up	9.48E-13	2.35E-12	AAEL003789-FA OS=Aedes aegypti GN=AAEL003789 PE=4 SV=1
CCG003459.1	35726	1771	4453	2.13008453	4.357308584	1.032526604	Up	0	0	AAEL013189-FA OS=Aedes aegypti GN=AAEL013189 PE=4 SV=1
CCG007457.1	2226	323	4456	6.23505134	69.97937165	3.488456366	Up	1.55E-13	4.61E-13	AAEL007544-FA OS=Aedes aegypti GN=AAEL007544 PE=4 SV=1
CCG008958.1	2076	823	4466	17.0347262	75.2040771	2.142332119	Up	7.96E-13	2.00E-12	AAEL009326-FA OS=Aedes aegypti GN=AAEL009326 PE=4 SV=1
CCG021947.1	3323	705	4470	9.11635958	47.02482611	2.366892873	Up	7.66E-13	1.92E-12	AAEL002798-FA OS=Aedes aegypti GN=AAEL002798 PE=4 SV=1
CCG011635.1	2847	11301	4477	170.565849	54.97302603	-1.633533012	Down	0	0	AAEL004805-FA OS=Aedes aegypti GN=AAEL004805 PE=4 SV=1
CCG022154.1	12321	12881	4480	44.9227458	12.7110689	-1.821360758	Down	0	0	AAEL013559-FA OS=Aedes aegypti GN=AAEL013559 PE=4 SV=1
CCG025914.2	1254	15663	4496	536.710489	125.336852	-2.09833342	Down	0	0	AAEL005170-FA OS=Aedes aegypti GN=AAEL005170 PE=4 SV=1
CCG013094.2	5811	774	4498	5.72338231	27.05942676	2.241191393	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_16871 PE=4 SV=1
CCG024993.1	11511	1694	4508	6.32358009	13.69054919	1.114366845	Up	6.33E-13	1.61E-12	AAEL002628-FA OS=Aedes aegypti GN=AAEL002628 PE=4 SV=1
CCG002966.1	561	748	4516	57.2929795	281.4110123	2.296248515	Up	4.45E-13	1.17E-12	60S ribosomal protein L17 OS=Culex quinquefasciatus GN=CpipJ_CPIJ015624 PE=3 SV=1

CCG010940.1	3243	1307	4516	17.3177438	48.68072091	1.491099549	Up	3.63E-12	8.42E-12	AAEL002986-PA OS=Aedes aegypti GN=AAEL002986 PE=4 SV=1
CCG006287.1	837	872	4517	44.7665574	188.6577493	2.075278077	Up	4.80E-13	1.25E-12	AAEL011542-PA OS=Aedes aegypti GN=AAEL011542 PE=3 SV=1
CCG020111.1	745	11	4521	0.63445246	212.1427776	8.385307787	Up	2.67E-07	5.03E-07	-
CCG019787.1	6215	917	4527	6.34002359	25.46357505	2.005874871	Up	0	0	AAEL000430-PA OS=Aedes aegypti GN=AAEL000430 PE=4 SV=1
CCG014045.1	2945	1561	4532	22.776148	53.79657397	1.239990528	Up	5.93E-12	1.34E-11	AAEL012217-PA OS=Aedes aegypti GN=AAEL012217 PE=4 SV=1
CCG017125.1	1833	1481	4541	34.7180453	86.60421976	1.3187516	Up	5.54E-12	1.25E-11	AAEL012845-PA OS=Aedes aegypti GN=AAEL012845 PE=4 SV=1
CCG000217.1	2246	896	4545	17.1419778	70.74148171	2.045022861	Up	6.75E-14	2.21E-13	Ribonucleotide reductase R2 subunit OS=Aedes albopictus PE=2 SV=1
CCG025514.1	1892	384	4548	8.72113007	84.03289786	3.268367242	Up	1.03E-12	2.55E-12	AAEL009882-PA OS=Aedes aegypti GN=AAEL009882 PE=4 SV=1
CCG025948.4	18727	841	4554	1.92970293	8.501094271	2.13926979	Up	1.23E-12	3.01E-12	Glucosamine-6-phosphate isomerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ008074 PE=3 SV=1
CCG019902.2	4451	710	4561	6.854305	35.82221981	2.385772444	Up	9.64E-14	3.03E-13	AAEL002369-PA OS=Aedes aegypti GN=AAEL002369 PE=4 SV=1
CCG005123.1	882	1496	4566	72.8829059	180.9744804	1.312133888	Up	1.42E-12	3.44E-12	AAEL003990-PA OS=Aedes aegypti GN=AAEL003990 PE=4 SV=1
CCG000669.1	2694	1092	4569	17.4175762	59.28892597	1.76721879	Up	3.67E-12	8.53E-12	AAEL009105-PA OS=Aedes aegypti GN=AAEL009105 PE=4 SV=1
CCG003625.1	1970	1623	4578	35.4009539	81.23806653	1.198367667	Up	4.27E-12	9.85E-12	AAEL003285-PA OS=Aedes aegypti GN=AAEL003285 PE=2 SV=1
CCG002038.1	1224	20972	4578	736.242871	130.7508097	-2.493361879	Down	0	0	ATP synthase subunit gamma OS=Aedes aegypti GN=AAEL008848 PE=3 SV=1
CCG007944.1	1224	11657	4579	409.230553	130.7793704	-1.645778876	Down	0	0	Arginine kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ007538 PE=3 SV=1

CCG010717.1	2610	124	4581	2.04147398	61.35780302	4.909563741	Up	1.61E-13	4.77E-13	AAEL014683-PA OS=Reedes aegypti GN=AAEL014683 PE=4 SV=1
CCG003149.1	1494	677	4590	19.4715598	107.4019347	2.463579619	Up	4.03E-13	1.07E-12	AAEL013950-PA OS=Reedes aegypti GN=AAEL013950 PE=3 SV=1
CCG021221.1	1869	1283	4608	29.4971479	86.18926665	1.54693275	Up	9.34E-13	2.32E-12	AAEL001007-PA OS=Reedes aegypti GN=AAEL001007 PE=4 SV=1
CCG023452.1	3210	1336	4608	17.8839768	50.18309638	1.488533913	Up	0	0	AAEL005455-PA OS=Reedes aegypti GN=AAEL005455 PE=4 SV=1
CCG025472.1	3117	354	4612	4.8801046	51.72523981	3.405884451	Up	0	0	AAEL004228-PA OS=Reedes aegypti GN=AAEL004228 PE=4 SV=1
CCG006884.1	2643	414	4614	6.73078703	61.02818351	3.180628535	Up	8.70E-13	2.17E-12	Phosphatase zC gamma OS=Culex quinquefasciatus GN=CpipJ_CPIJ017206 PE=3 SV=1
CCG027273.1	1654	1228	4615	31.9025599	97.5407783	1.612333291	Up	4.21E-12	9.72E-12	AAEL001188-PA OS=Reedes aegypti GN=AAEL001188 PE=4 SV=1
CCG015076.1	2561	643	4616	10.7885745	63.00952971	2.546065785	Up	0	0	AAEL005736-PA OS=Reedes aegypti GN=AAEL005736 PE=4 SV=1
CCG014707.1	504	149	4624	12.7033541	320.7283171	4.658070366	Up	2.33E-13	6.63E-13	-
CCG026971.1	4626	458	4624	4.25424523	34.94316295	3.038035098	Up	1.14E-12	2.79E-12	AAEL008730-PA OS=Reedes aegypti GN=AAEL008730 PE=4 SV=1
CCG018120.1	2901	1554	4626	23.0179137	55.74525624	1.276091966	Up	4.78E-12	1.09E-11	AAEL001448-PA OS=Reedes aegypti GN=AAEL001448 PE=4 SV=1
CCG016323.1	65605	1597	4629	1.0459975	2.466608691	1.237649454	Up	6.60E-12	1.48E-11	AAEL013859-PA OS=Reedes aegypti GN=AAEL013859 PE=4 SV=1
CCG002819.1	2439	43	4637	0.757564	66.46229167	6.45502446	Up	8.46E-14	2.70E-13	AAEL003261-PA OS=Reedes aegypti GN=AAEL003261 PE=4 SV=1
CCG017917.1	4192	1241	4643	12.7207635	38.71929367	1.605867369	Up	3.14E-12	7.34E-12	AAEL013866-PA OS=Reedes aegypti GN=AAEL013866 PE=4 SV=1
CCG028242.1	1122	1702	4648	65.1822534	144.8182446	1.151692236	Up	5.16E-12	1.17E-11	AAEL004589-PA OS=Reedes aegypti GN=AAEL004589 PE=3 SV=1
CCG016955.1	1215	1207	4665	42.6868063	134.2225195	1.652764609	Up	6.35E-12	1.43E-11	AAEL004501-PA OS=Reedes aegypti GN=AAEL004501 PE=3 SV=1
CCG021215.1	2472	353	4672	6.13605029	66.07001176	3.428613389	Up	0	0	AAEL004818-PA OS=Reedes aegypti GN=AAEL004818 PE=4 SV=1

CCG005355.1	6053	826	4674	5.86370408	26.99405016	2.202757251	Up	1.49E-13	4.45E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013516 PE=4 SV=1
CCG003693.1	35225	1525	4680	1.86029369	4.644563103	1.320012491	Up	7.80E-12	1.75E-11	AAEL005080-PA OS=Aedes aegypti GN=AAEL005080 PE=4 SV=1
CCG027568.1	3002	490	4681	7.01371418	54.51022437	2.958276315	Up	0	0	AAEL004870-PA OS=Aedes aegypti GN=CYP18A1 PE=3 SV=1
CCG016182.1	1458	11537	4685	340.014971	112.331637	-1.597833968	Down	0	0	AGAP001419-PC OS=Anopheles gambiae GN=AgaP_AGAP001419 PE=4 SV=1
CCG027465.1	3086	592	4687	8.24305991	53.09444045	2.687308918	Up	4.13E-13	1.10E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012359 PE=4 SV=1
CCG006534.1	2670	84	4692	1.35185682	61.43229761	5.505983079	Up	0	0	AAEL008098-PA OS=Aedes aegypti GN=PIWI2 PE=4 SV=1
CCG005401.1	8238	1602	4702	8.35609551	19.9531218	1.255713595	Up	1.61E-12	3.87E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003108 PE=4 SV=1
CCG003339.1	2259	12109	4706	230.332234	72.8258745	-1.661193298	Down	0	0	AAEL012160-PA OS=Aedes aegypti GN=AAEL012160 PE=4 SV=1
CCG008539.2	2247	1435	4708	27.4417308	73.24591324	1.416376787	Up	5.38E-12	1.22E-11	T-complex protein 1 subunit alpha OS=Aedes aegypti GN=AAEL009320 PE=3 SV=1
CCG008734.1	3174	8034	4710	108.764602	51.87570372	-1.068078198	Down	0	0	AAEL007979-PA OS=Aedes aegypti GN=AAEL007979 PE=4 SV=1
CCG002170.1	1560	57	4711	1.570048	105.569514	6.071240806	Up	0	0	AAEL002554-PA OS=Aedes aegypti GN=AAEL002554 PE=4 SV=1
CCG022824.1	13389	950	4716	3.04886458	12.31333432	2.013877504	Up	7.21E-13	1.81E-12	AAEL012187-PA OS=Aedes aegypti GN=AAEL012187 PE=4 SV=1
CCG006600.1	3559	930	4720	11.2283937	46.36219903	2.045797442	Up	0	0	AAEL008319-PA OS=Aedes aegypti GN=AAEL008319 PE=4 SV=1
CCG018476.1	660	9178	4728	597.539734	250.4283827	-1.254636567	Down	0	0	AAEL007162-PA OS=Aedes aegypti GN=AAEL007162 PE=3 SV=1
CCG005050.1	3137	1430	4736	19.5877337	52.77730277	1.429967138	Up	4.48E-12	1.03E-11	AAEL003367-PA OS=Aedes aegypti GN=AAEL003367 PE=4 SV=1

CCG018418.2	3737	1009	4739	11.6019433	44.33162259	1.93396969	Up	2.35E-12	5.55E-12	AAEL000427-PA OS=Aedes aegypti GN=AAEL000427 PE=4 SV=1
CCG022020.1	11047	1839	4746	7.15319471	15.01873645	1.070103825	Up	0	0	AAEL011584-PA OS=Aedes aegypti GN=AAEL011584 PE=3 SV=1
CCG009084.2	3873	1146	4755	12.7145148	42.91934058	1.755151501	Up	4.08E-12	9.43E-12	S-hydroxyisobutyrate dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ000400 PE=4 SV=1
CCG003712.1	22503	15587	4755	29.7635539	7.386864243	-2.010512832	Down	0	0	AAEL002436-PA (Fragment) OS=Aedes aegypti GN=AAEL002436 PE=2 SV=1
CCG026842.1	1699	1757	4760	44.4366237	97.94078719	1.140160586	Up	0	0	AAEL003676-PA OS=Aedes aegypti GN=AAEL003676 PE=4 SV=1
CCG010262.1	3414	1324	4763	16.6643025	48.77160875	1.54928063	Up	1.05E-12	2.59E-12	AGAP009189-PA (Fragment) OS=Anopheles gambiae GN=AGAP009189 PE=4 SV=4
CCG028049.1	1539	1435	4764	40.0659969	108.213925	1.43343588	Up	5.38E-12	1.22E-11	AAEL002866-PA OS=Aedes aegypti GN=AAEL002866 PE=3 SV=1
CCG018650.1	4569	404	4768	3.79946876	36.48086313	3.263270242	Up	1.18E-13	3.65E-13	AAEL005597-PA OS=Aedes aegypti GN=AAEL005597 PE=4 SV=1
CCG004132.1	5895	1335	4769	9.73105949	28.28091975	1.539160245	Up	6.03E-12	1.36E-11	AAEL003487-PA (Fragment) OS=Aedes aegypti GN=AAEL003487 PE=4 SV=1
CCG010244.1	10200	1273	4774	5.36279139	16.36184444	1.609279351	Up	4.95E-12	1.13E-11	AAEL007069-PA OS=Aedes aegypti GN=AAEL007069 PE=4 SV=1
CCG005319.1	9360	12630	4785	57.981597	17.87129854	-1.697950605	Down	0	0	AAEL003078-PA OS=Aedes aegypti GN=AAEL003078 PE=4 SV=1
CCG006745.1	2399	23006	4787	412.072411	69.75626132	-2.562503252	Down	0	0	AAEL002764-PA OS=Aedes aegypti GN=AAEL002764 PE=3 SV=1
CCG012844.1	9873	934	4796	4.0649987	16.98165658	2.062650408	Up	3.46E-13	9.34E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000695 PE=4 SV=1
CCG000294.1	3424	1349	4812	16.9293727	49.12944738	1.537059498	Up	8.73E-12	1.95E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_02441 PE=3 SV=1

CCG015426.1	6615	498	4813	3.23490973	25.43525111	2.97503198	Up	3.42E-13	9.25E-13	AAEL004375-PA OS=Aedes aegypti GN=AAEL004375 PE=4 SV=1
CCG017215.1	3156	785	4813	10.6879726	53.31247975	2.318485068	Up	0	0	AAEL001440-PA OS=Aedes aegypti GN=AAEL001440 PE=4 SV=1
CCG013216.1	2470	15316	4829	266.447148	68.34555406	-1.962929993	Down	0	0	AAEL003740-PA OS=Aedes aegypti GN=AAEL003746 PE=4 SV=1
CCG009712.1	4551	1439	4843	13.5867827	37.20126003	1.453147615	Up	3.28E-12	7.67E-12	AAEL002345-PA OS=Aedes aegypti GN=AAEL002345 PE=4 SV=1
CCG004388.1	3507	8532	4854	104.538858	48.38536511	-1.111396653	Down	0	0	Pyruvate carboxylase OS=Aedes aegypti PE=2 SV=1
CCG025919.1	3888	469	4858	5.18333476	43.67986331	3.075015866	Up	0	0	AAEL007823-PA OS=Aedes aegypti GN=AAEL007823 PE=4 SV=1
CCG010477.1	3206	1542	4862	20.6672897	53.01532803	1.359060332	Up	2.49E-12	5.89E-12	AAEL008216-PA OS=Aedes aegypti GN=AAEL008216 PE=4 SV=1
CCG017381.1	2911	1834	4871	27.0719661	58.49596914	1.111537548	Up	0	0	AAEL002804-PA OS=Aedes aegypti GN=AAEL002804 PE=4 SV=1
CCG007203.1	2129	13921	4887	280.968377	80.24476214	-1.807928636	Down	0	0	AAEL013431-PA (Fragment) OS=Aedes aegypti GN=AAEL013431 PE=4 SV=1
CCG005131.1	2481	24886	4887	431.01363	68.85977371	-2.646000147	Down	0	0	AAEL006323-PA OS=Aedes aegypti GN=AAEL006323 PE=4 SV=1
CCG003104.1	3153	1835	4891	25.0077586	54.22801513	1.116662605	Up	0	0	AAEL002152-PA OS=Aedes aegypti GN=AAEL002152 PE=4 SV=1
CCG003603.1	11322	855	4892	3.24493226	15.10474209	2.218741283	Up	0	0	AAEL013278-PA OS=Aedes aegypti GN=AAEL013278 PE=4 SV=1
CCG010025.1	3126	682	4895	9.37471497	54.74112758	2.545778419	Up	1.08E-12	2.66E-12	AAEL003359-PA (Fragment) OS=Aedes aegypti GN=AAEL003359 PE=4 SV=1
CCG003146.1	3296	622	4906	8.10897298	52.03437677	2.681873952	Up	0	0	AAEL013275-PA OS=Aedes aegypti GN=AAEL013275 PE=4 SV=1
CCG002418.1	1533	1928	4909	54.0415188	111.9440187	1.050637319	Up	0	0	AAEL002023-PA OS=Aedes aegypti GN=AAEL002023 PE=3 SV=1
CCG023790.1	23208	1244	4913	2.30327257	7.400465951	1.683930957	Up	1.32E-12	3.20E-12	AAEL002729-PA OS=Aedes aegypti GN=AAEL002729 PE=4 SV=1

CCG018231.1	2061	105	4914	2.18914223	83.35030182	5.250749829	Up	0	0	AAEL001666-PA OS=Aedes aegypti GN=AAEL001666 PE=4 SV=1
CCG006265.1	1456	1794	4941	52.9448516	118.6324489	1.163936369	Up	0	0	Methionine aminopeptidase OS=Aedes aegypti GN=AAEL004738 PE=3 SV=1
CCG009035.1	1455	451	4942	13.3191411	118.7380095	3.156208876	Up	1.51E-12	3.65E-12	AAEL000672-PA OS=Aedes aegypti GN=AAEL000672 PE=3 SV=1
CCG008035.1	1698	1048	4942	26.5207785	101.7454675	1.939769498	Up	3.41E-12	7.95E-12	Tripe functional domain OS=Culex quinquefasciatus GN=CpipJ_CPIJ002249 PE=4 SV=1
CCG023748.1	7032	698	4943	4.26519834	24.5732028	2.526401169	Up	0	0	AAEL009265-PA OS=Aedes aegypti GN=AAEL009265 PE=4 SV=1
CCG025300.1	1605	1245	4957	33.3316633	107.9677121	1.69563472	Up	4.85E-12	1.11E-11	AAEL006389-PA (Fragment) OS=Aedes aegypti GN=AAEL006389 PE=3 SV=1
CCG017883.2	5333	1485	4959	11.9651333	32.50667439	1.441899498	Up	0	0	AAEL005557-PA OS=Aedes aegypti GN=AAEL005557 PE=4 SV=1
CCG002465.1	2126	1566	4961	31.6512721	81.57479354	1.365859948	Up	3.46E-13	9.36E-13	AAEL010063-PA OS=Aedes aegypti GN=AAEL010063 PE=3 SV=1
CCG001172.1	5071	518	4965	4.38933594	34.2275378	2.963082919	Up	0	0	AAEL009392-PA OS=Aedes aegypti GN=AAEL009392 PE=4 SV=1
CCG027635.1	3780	302	4979	3.43303171	46.04689419	3.745548765	Up	4.59E-13	1.20E-12	AAEL003382-PA OS=Aedes aegypti GN=AAEL003382 PE=4 SV=1
CCG002823.1	714	1112	4983	66.9220517	243.9735198	1.86617099	Up	5.39E-12	1.22E-11	AAEL004284-PA OS=Aedes aegypti GN=AAEL004284 PE=4 SV=1
CCG005015.1	231	35007	4983	6511.86796	754.0999704	-3.110243749	Down	0	0	AAEL014039-PA OS=Aedes aegypti GN=AAEL014039 PE=4 SV=1
CCG019032.1	2703	1250	4991	19.8713164	64.54930054	1.699714013	Up	5.51E-13	1.42E-12	AAEL010599-PA OS=Aedes aegypti GN=AAEL010599 PE=4 SV=1
CCG014378.1	1500	921	4993	26.3834171	116.3644506	2.140947049	Up	0	0	AAEL011314-PA OS=Aedes aegypti GN=AAEL011314 PE=2 SV=1
CCG002723.1	3909	488	4995	5.36434651	44.67039971	3.057844829	Up	9.43E-13	2.34E-12	AAEL014733-PA OS=Aedes aegypti GN=AAEL014733 PE=4 SV=1

CCG010190.1	1539	745	5003	20.8008137	113.6427932	2.449794325	Up	0	0	O-acyltransferase (Fragment) OS=Aedes aegypti GN=AAEL001204 PE=3 SV=1
CCG015641.1	3954	1102	5013	11.9758846	44.32115363	1.887863214	Up	3.03E-12	7.09E-12	AAEL012607-PA OS=Aedes aegypti GN=AAEL012607 PE=4 SV=1
CCG001217.1	3708	1325	5017	15.3546112	47.29926499	1.623145782	Up	1.87E-12	4.48E-12	Ubiquitin carboxyl-terminal hydrolase OS=Aedes aegypti GN=AAEL004980 PE=3 SV=1
CCG001811.1	4932	800	5022	6.96994884	35.59620153	2.352503327	Up	9.48E-13	2.35E-12	AAEL005801-PA OS=Aedes aegypti GN=AAEL005801 PE=3 SV=1
CCG022370.1	3321	1309	5025	16.9368812	52.8953149	1.642971703	Up	2.04E-12	4.87E-12	AAEL005320-PA (Fragment) OS=Aedes aegypti GN=AAEL005320 PE=4 SV=1
CCG002373.1	2755	207	5031	3.22857897	63.83850833	4.305455721	Up	6.88E-15	2.47E-14	AAEL014051-PA OS=Aedes aegypti GN=AAEL014051 PE=4 SV=1
CCG010564.2	1167	1611	5033	59.3181169	150.7669297	1.345775308	Up	0	0	Importin subunit alpha OS=Aedes aegypti GN=AAEL011657 PE=3 SV=1
CCG025081.1	1375	10	5034	0.31250716	127.9854293	8.67787461	Up	5.43E-07	1.01E-06	AAEL001401-PA OS=Aedes aegypti GN=AAEL001401 PE=4 SV=1
CCG020033.1	4326	460	5034	4.56913498	40.67960362	3.154312654	Up	0	0	Condensin complex subunit 1 OS=Aedes aegypti GN=AAEL004634 PE=3 SV=1
CCG005153.1	8153	745	5038	3.92646293	21.60183961	2.459851995	Up	0	0	histone-lysine N- methyltransferase ash1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001202 PE=4 SV=1
CCG027231.1	6186	1002	5040	6.9601801	28.48201017	2.032854429	Up	3.07E-12	7.20E-12	AAEL005712-PA OS=Aedes aegypti GN=AAEL005712 PE=4 SV=1
CCG020413.1	4217	1159	5040	11.809799	41.78081929	1.822856371	Up	3.95E-12	9.12E-12	AAEL002581-PA OS=Aedes aegypti GN=AAEL002581 PE=4 SV=1
CCG003143.1	2010	555	5044	11.8647775	87.7261433	2.886321803	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014888 PE=4 SV=1
CCG006337.1	21929	347	5051	0.67994427	8.052088831	3.565874678	Up	0	0	AAEL007698-PA OS=Aedes aegypti GN=PIWI4 PE=4 SV=1

CCG015097.3	5165	1132	5051	9.41756817	34.18669041	1.860008287	Up	5.10E-12	1.16E-11	AAEL000212-PA OS=Aedes aegypti GN=AAEL000212 PE=4 SV=1
CCG001435.1	2124	2054	5053	41.5535946	83.16580628	1.001017202	Up	0	0	AAEL010991-PA OS=Aedes aegypti GN=AAEL010991 PE=4 SV=1
CCG009699.2	3216	1254	5054	16.7549898	54.93754067	1.71320152	Up	1.97E-12	4.69E-12	AAEL003939-PA OS=Aedes aegypti GN=AAEL003939 PE=4 SV=1
CCG013833.1	609	431	5079	30.4104362	291.5485841	3.261097903	Up	0	0	AAEL002372-PB OS=Aedes aegypti GN=RpS11 PE=2 SV=1
CCG008130.1	2770	20920	5081	324.522328	64.12382826	-2.339385279	Down	0	0	AAEL004221-PA OS=Aedes aegypti GN=AAEL004221 PE=4 SV=1
CCG027463.1	2944	201	5084	2.93373528	60.36952415	4.363009828	Up	0	0	AAEL011838-PA OS=Aedes aegypti GN=AAEL011838 PE=4 SV=1
CCG019245.2	6177	1548	5084	10.7685202	28.77252373	1.417871763	Up	3.12E-12	7.30E-12	AGAP003873-PA OS=Anopheles gambiae GN=AgaP_AGAP003873 PE=4 SV=5
CCG026313.1	3574	444	5085	5.33815394	49.73778326	3.219929396	Up	6.43E-13	1.64E-12	AAEL007201-PA (Fragment) OS=Aedes aegypti GN=AAEL007201 PE=4 SV=1
CCG002282.1	2116	1661	5086	33.7300232	84.02542328	1.316792593	Up	6.26E-12	1.41E-11	AAEL004300-PA OS=Aedes aegypti GN=AAEL004300 PE=4 SV=1
CCG021083.1	5649	1569	5087	11.9347696	31.48039546	1.399282946	Up	0	0	AAEL004300-PA OS=Aedes aegypti GN=AAEL004300 PE=4 SV=1 DNA-directed RNA polymerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ016666 PE=3 SV=1
CCG023599.1	6642	1003	5096	6.48880515	26.82134574	2.047356875	Up	3.57E-12	8.30E-12	AAEL002709-PA OS=Aedes aegypti GN=AAEL002709 PE=4 SV=1
CCG014997.1	3111	110	5099	1.51934131	57.29741346	5.236952113	Up	0	0	AAEL002036-PA OS=Aedes aegypti GN=AAEL002036 PE=4 SV=1
CCG011971.1	2600	399	5101	6.59420158	68.58544993	3.378632653	Up	0	0	AAEL003071-PA OS=Aedes aegypti GN=AAEL003071 PE=4 SV=1
CCG021239.1	1248	520	5114	17.9040561	143.2505027	3.000181836	Up	1.17E-12	2.86E-12	AAEL003443-PA OS=Aedes aegypti GN=AAEL003443 PE=4 SV=1
CCG014780.2	2769	854	5119	13.2524931	64.62673124	2.285867235	Up	1.02E-12	2.51E-12	AAEL004946-PA (Fragment) OS=Aedes aegypti GN=AAEL004946 PE=4 SV=1

CCG018850.1	1683	3	5120	0.07659489	106.3496002	10.4392788	Up	5.28E-05	8.85E-05	AAEL000335-PA OS=Aedes aegypti GN=AAEL000335 PE=4 SV=1
CCG006613.1	6615	1266	5124	8.22368617	27.07879217	1.719306275	Up	2.71E-12	6.37E-12	AAEL010835-PA (Fragment) OS=Aedes aegypti GN=AAEL010835 PE=4 SV=1
CCG016109.1	1545	241	5135	6.70272236	116.1881885	4.115572429	Up	2.37E-13	6.72E-13	AAEL014365-PA (Fragment) OS=Aedes aegypti GN=AAEL014365 PE=4 SV=1
CCG026105.1	2586	731	5141	12.1465104	69.49748681	2.516418905	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003815 PE=4 SV=1
CCG008301.1	2383	1056	5143	19.041561	75.44709083	1.986313523	Up	2.35E-12	5.56E-12	AAEL010641-PA OS=Aedes aegypti GN=AAEL010641 PE=4 SV=1
CCG002713.1	1770	226	5146	5.48653109	101.6357583	4.211369983	Up	2.04E-14	7.16E-14	AAEL008863-PA OS=Aedes aegypti GN=AAEL008863 PE=4 SV=1
CCG001566.1	1440	843	5151	25.1551988	125.0486692	2.313561208	Up	2.68E-12	6.30E-12	AAEL011985-PA OS=Aedes aegypti GN=AAEL011985 PE=4 SV=1
CCG015053.1	828	1609	5153	83.5003659	217.5603867	1.38156147	Up	5.36E-12	1.22E-11	AAEL013775-PA OS=Aedes aegypti GN=AAEL013775 PE=4 SV=1
CCG016476.1	543	0	5157	0.001	332.0070595	18.34085439	Up	4.44E-16	1.63E-15	AAEL013262-PA OS=Aedes aegypti GN=AAEL013262 PE=4 SV=1
CCG011662.1	4685	1056	5164	9.68538735	38.53245278	1.99219237	Up	2.35E-12	5.56E-12	AAEL011534-PA OS=Aedes aegypti GN=AAEL011534 PE=4 SV=1
CCG007304.2	38717	721	5168	0.80019574	4.6662803	2.543848109	Up	0	0	AAEL012039-PA OS=Aedes aegypti GN=AAEL012039 PE=4 SV=1
CCG002947.1	5415	17338	5173	137.582504	33.39596782	-2.042551188	Down	0	0	Histone H3 OS=Gecko japonicus PE=2 SV=1
CCG000431.1	2229	21520	5180	414.853606	81.23996127	-2.352340871	Down	0	0	Glucose-6-phosphate isomerase OS=Aedes aegypti GN=AAEL012994 PE=3 SV=1
CCG011703.1	3777	1983	5181	22.5599639	47.95309292	1.08785911	Up	0	0	AAEL012753-PA (Fragment) OS=Aedes aegypti GN=AAEL012753 PE=4 SV=1
CCG020159.1	11061	885	5194	3.43804494	16.41563055	2.255409858	Up	0	0	AAEL013275-PA OS=Aedes aegypti GN=AAEL013275 PE=4 SV=1

CCG020056.1	61404	1997	5199	1.39747508	2.959873639	1.082713026	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008269 PE=4 SV=1
CCG016099.1	4960	1357	5204	11.7560342	36.67799845	1.641513445	Up	3.94E-12	9.11E-12	Spastin OS=Culex quinquefasciatus GN=CpipJ_CPIJ012667 PE=2 SV=1
CCG025030.1	804	232	5205	12.3992269	226.3157097	4.190014657	Up	2.91E-13	8.06E-13	AAEL003389-PA OS=Aedes aegypti GN=ATT PE=4 SV=1
CCG015096.1	9027	1030	5211	4.90293859	20.18030135	2.041229121	Up	4.77E-12	1.09E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000695 PE=4 SV=1
CCG011550.1	4131	341	5221	3.54700545	44.18231978	3.638795718	Up	2.98E-13	8.25E-13	AAEL012336-PA OS=Aedes aegypti GN=AAEL012336 PE=4 SV=1
CCG011179.1	1515	862	5221	24.4487863	120.4733749	2.300879588	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011773 PE=4 SV=1
CCG005040.1	4962	748	5228	6.47750131	36.83229967	2.50746217	Up	4.45E-13	1.17E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005688 PE=4 SV=1
CCG012522.1	2493	2049	5228	35.3168818	73.31001643	1.053652361	Up	0	0	AAEL002621-PA OS=Aedes aegypti GN=AAEL002621 PE=4 SV=1
CCG023316.1	17029	1839	5232	4.64039826	10.74060157	1.210754265	Up	0	0	Saccharopine dehydrogenase domain-containing protein OS=Aedes aegypti PE=2 SV=1
CCG016702.1	1738	704	5235	17.4054621	105.2972261	2.596855407	Up	1.23E-12	3.00E-12	AAEL009220-PA OS=Aedes aegypti GN=AAEL009220 PE=4 SV=1
CCG012377.1	3585	1282	5235	15.366025	51.04786022	1.732106479	Up	3.23E-12	7.55E-12	PTEN transcript variant 6 OS=Aedes aegypti PE=2 SV=1
CCG006625.1	5045	462	5237	3.93498858	36.28870078	3.205089052	Up	0	0	AAEL009392-PA OS=Aedes aegypti GN=AAEL009392 PE=4 SV=1
CCG010008.1	11493	13587	5238	50.7987283	15.93243311	-1.672825777	Down	0	0	AGAP009638-PA (Fragment) OS=Anopheles gambiae GN=AGAP009638 PE=4 SV=4
CCG016017.1	4325	2119	5241	21.0526862	42.36215689	1.008771725	Up	0	0	Tyrosine-protein kinase shark OS=Culex quinquefasciatus GN=CpipJ_CPIJ011800 PE=4 SV=1

CCG000358.1	2076	203	5274	4.20176114	88.81018868	4.401659142	Up	9.28E-14	2.93E-13	AAEL007813-PA OS=Aedes aegypti GN=AAEL007813 PE=4 SV=1
CCG009822.1	6621	1246	5279	8.08643548	27.87263904	1.785273801	Up	2.10E-12	5.00E-12	AAEL007586-PA OS=Aedes aegypti GN=AAEL007586 PE=4 SV=1
CCG000112.1	360	1669	5280	199.212464	512.7213927	1.363867179	Up	3.31E-12	7.71E-12	AAEL000068-PA OS=Aedes aegypti GN=AAEL000068 PE=4 SV=1
CCG012770.1	4590	11125	5283	104.147777	40.23629111	-1.372062817	Down	0	0	AAEL010382-PA OS=Aedes aegypti GN=AAEL010382 PE=4 SV=1
CCG025614.1	4377	11406	5288	111.974593	42.23426264	-1.406685652	Down	0	0	Ckz1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008998 PE=4 SV=1
CCG023890.1	1551	1611	5303	44.6320067	119.5253009	1.421165458	Up	0	0	AAEL002023-PA OS=Aedes aegypti GN=AAEL002023 PE=3 SV=1
CCG027398.1	4977	1928	5306	16.6456999	37.26916145	1.162832828	Up	0	0	Steroid regulatory element-binding protein cleavage-activating protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ013729 PE=4 SV=1
CCG024115.1	1488	1862	5307	53.7699233	124.6798218	1.21335668	Up	0	0	AAEL003634-PA OS=Aedes aegypti GN=AAEL003634 PE=2 SV=1
CCG004346.1	3680	508	5323	5.93169163	50.56600741	3.091652368	Up	6.20E-13	1.58E-12	AAEL010444-PA (Fragment) OS=Aedes aegypti GN=AAEL010444 PE=4 SV=1
CCG014654.1	2184	547	5332	10.7621084	85.3468552	2.987377246	Up	0	0	AAEL013905-PB (Fragment) OS=Aedes aegypti GN=AAEL013905 PE=4 SV=1
CCG027745.1	13689	12815	5335	40.2262509	13.62425353	-1.561960099	Down	0	0	AAEL009847-PA OS=Aedes aegypti GN=AAEL009847 PE=4 SV=1
CCG011536.1	3932	530	5341	5.79195304	47.48528897	3.035358823	Up	0	0	AAEL001099-PA OS=Aedes aegypti GN=AAEL001099 PE=4 SV=1
CCG013339.1	17750	49087	5368	118.831288	10.57217069	-3.490571217	Down	0	0	Fructose-bisphosphate aldolase OS=Aedes aegypti GN=AAEL005766 PE=3 SV=1
CCG027495.1	9020	12535	5369	59.7145924	20.80831352	-1.520923482	Down	0	0	AGAP003857-PA OS=Anopheles gambiae GN=AgaP_AGAP003857 PE=3 SV=5

CCG018028.1	4049	21399	5373	227.095419	46.3894347	-2.291430432	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000251 PE=4 SV=1
CCG023263.1	2477	621	5374	10.7727918	75.84407726	2.815644352	Up	0	0	AAEL007158-PA OS=Aedes aegypti GN=AAEL007158 PE=4 SV=1
CCG021632.2	5619	39455	5375	301.721103	33.44024518	-3.173558276	Down	0	0	EF1r domain-binding protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ012054 PE=4 SV=1
CCG002698.1	3615	375	5386	4.45744135	52.08444777	3.546564933	Up	0	0	TSC1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000640 PE=4 SV=1
CCG019725.1	971	32	5392	1.41609836	194.124643	7.098917985	Up	1.77E-13	5.16E-13	AAEL013338-PA OS=Aedes aegypti GN=AAEL013338 PE=3 SV=1
CCG027793.1	1743	2073	5409	51.1051405	108.4849794	1.085954985	Up	0	0	AAEL003285-PA OS=Aedes aegypti GN=AAEL003285 PE=2 SV=1
CCG015622.2	4028	549	5413	5.85659988	46.97843897	3.003863538	Up	2.32E-13	6.62E-13	AAEL007863-PA OS=Aedes aegypti GN=AAEL007863 PE=4 SV=1
CCG000959.2	3828	1536	5414	17.2417744	49.44203512	1.519829875	Up	2.61E-12	6.14E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_15687 PE=4 SV=1
CCG015169.1	3296	1654	5415	21.5630889	57.43296988	1.413315307	Up	2.93E-12	6.87E-12	AAEL003415-PA OS=Aedes aegypti GN=AAEL003415 PE=4 SV=1
CCG000936.1	1488	708	5440	20.445277	127.8044527	2.64409859	Up	9.24E-13	2.30E-12	AAEL008578-PA OS=Aedes aegypti GN=AAEL008578 PE=4 SV=1
CCG018534.1	942	1927	5441	87.9009327	201.9193035	1.199828461	Up	0	0	AAEL002160-PA OS=Aedes aegypti GN=AAEL002160 PE=4 SV=1
CCG028185.1	1411	1360	5442	41.4166117	134.8284495	1.702843509	Up	1.23E-12	3.01E-12	Insulin-like peptide 6 splice variant A OS=Aedes aegypti PE=2 SV=1
CCG016240.1	3769	579	5445	6.60108154	50.5035333	2.935609999	Up	7.33E-13	1.84E-12	AAEL004348-PA OS=Aedes aegypti GN=AAEL004348 PE=4 SV=1
CCG024762.1	3161	11498	5448	156.300541	60.25077249	-1.375271124	Down	0	0	AAEL006446-PA OS=Aedes aegypti GN=AAEL006446 PE=4 SV=1
CCG004456.1	4435	707	5452	6.84996671	42.97463922	2.649316646	Up	0	0	AAEL002369-PA OS=Aedes aegypti GN=AAEL002369 PE=4 SV=1

CCG025456.1	1434	1090	5468	32.6617927	133.2997611	2.028998312	Up	4.87E-12	1.11E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007053 PE=4 SV=1
CCG015235.1	35415	1194	5479	1.44870431	5.408341054	1.900422971	Up	1.96E-12	4.67E-12	AAEL009265-PA OS=Aedes aegypti GN=AAEL009265 PE=4 SV=1
CCG011740.1	843	14	5517	0.71361362	228.7838825	8.324626547	Up	3.07E-08	6.00E-08	Salivary antigen-5 related protein AG5-4 OS=Aedes albopictus PE=2 SV=1
CCG001970.1	1833	466	5523	10.9241115	105.3325492	3.269363471	Up	0	0	AAEL007799-PA OS=Aedes aegypti GN=AAEL007799 PE=4 SV=1
CCG021203.1	1762	918	5528	22.387183	109.6761373	2.292504761	Up	1.42E-12	3.43E-12	AAEL011892-PA OS=Aedes aegypti GN=AAEL011892 PE=4 SV=1
CCG028336.1	7578	958	5529	5.43217284	25.50597945	2.231234214	Up	1.10E-12	2.71E-12	AAEL000295-PA OS=Aedes aegypti GN=AAEL000295 PE=4 SV=1
CCG015998.1	1755	868	5531	21.2522676	110.1733498	2.374086598	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_03313 PE=4 SV=1
CCG022499.1	2466	531	5533	9.25260709	78.4363931	3.083591361	Up	2.40E-13	6.76E-13	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000842 PE=4 SV=1
CCG013110.1	1314	934	5554	30.5431751	147.7612399	2.274345926	Up	3.46E-13	9.35E-13	AAEL012312-PA OS=Aedes aegypti GN=AAEL012312 PE=4 SV=1
CCG001881.1	4115	410	5555	4.28131013	47.19154981	3.462404301	Up	1.25E-12	3.04E-12	AAEL002714-PA OS=Aedes aegypti GN=AAEL002714 PE=3 SV=1
CCG025171.1	2140	1416	5557	28.4323104	90.7771701	1.674798179	Up	1.35E-12	3.28E-12	AAEL005947-PB OS=Aedes aegypti GN=AAEL005947 PE=4 SV=1
CCG003699.1	9903	1914	5562	8.30496537	19.63424572	1.241326119	Up	0	0	Lysosomal trafficking regulator OS=Culex quinquefasciatus GN=CpipJ_CPIJ005368 PE=4 SV=1
CCG011072.1	2904	508	5575	7.51674421	67.11170558	3.158384607	Up	6.20E-13	1.58E-12	AAEL001891-PA OS=Aedes aegypti GN=AAEL001891 PE=4 SV=1
CCG004169.1	4983	1273	5575	10.9774177	39.11145756	1.83305259	Up	4.95E-12	1.13E-11	Conductor enhancer of ksr OS=Culex quinquefasciatus GN=CpipJ_CPIJ001765 PE=4 SV=1

CCG011096.1	5973	9488	5577	68.2566285	32.64060096	-1.064301535	Down	0	0	AAEL005175-PA OS=Aedes aegypti GN=AAEL005175 PE=4 SV=1
CCG017499.1	15195	1676	5584	4.73953769	12.84679286	1.438589996	Up	1.21E-12	2.96E-12	AAEL008015-PA OS=Aedes aegypti GN=AAEL008015 PE=4 SV=1
CCG019244.1	4559	1426	5595	13.4404127	42.90229404	1.674477353	Up	2.31E-12	5.48E-12	AGAP003872-PA OS=Anopheles gambiae GN=AgaP_AGAP003872 PE=4 SV=4
CCG021178.1	2370	917	5598	16.6258425	82.57233476	2.312231052	Up	0	0	AAEL002370-PA OS=Aedes aegypti GN=AAEL002370 PE=4 SV=1
CCG000436.1	3025	16104	5619	228.755242	64.93572138	-1.816720568	Down	0	0	AAEL008953-PA OS=Aedes aegypti GN=AAEL008953 PE=4 SV=1
CCG009103.1	1203	958	5625	34.2186249	163.4582767	2.256068739	Up	1.10E-12	2.71E-12	AAEL013408-PA OS=Aedes aegypti GN=AAEL013408 PE=4 SV=1
CCG017045.1	3093	632	5637	8.78010743	63.71154419	2.859244308	Up	0	0	AAEL008806-PA OS=Aedes aegypti GN=AAEL008806 PE=4 SV=1
CCG028581.1	1805	21312	5652	507.352346	109.4649198	-2.212519413	Down	0	0	AAEL011381-PA OS=Aedes aegypti GN=AAEL011381 PE=3 SV=1
CCG027266.1	3369	1120	5666	14.2849815	58.79299382	2.041145073	Up	3.68E-12	8.54E-12	AAEL001069-PA OS=Aedes aegypti GN=AAEL001069 PE=4 SV=1
CCG025794.2	1827	816	5688	19.191737	108.8356203	2.503593611	Up	2.12E-12	5.04E-12	Zinc finger protein 595 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008058 PE=4 SV=1
CCG022430.1	4482	20601	5694	197.505467	44.41151895	-2.152886767	Down	0	0	AAEL009838-PA OS=Aedes aegypti GN=AAEL009838 PE=4 SV=1
CCG008920.1	2661	548	5697	8.84908477	74.84302998	3.080267812	Up	9.78E-13	2.42E-12	AAEL012918-PA OS=Aedes aegypti GN=AAEL012918 PE=4 SV=1
CCG022974.1	1560	1019	5700	28.068051	127.7321651	2.186121072	Up	3.75E-12	8.69E-12	AAEL002527-PA OS=Aedes aegypti GN=AAEL002527 PE=4 SV=1
CCG024593.1	2478	2043	5708	35.4266214	80.5253607	1.184609335	Up	0	0	AAEL003812-PA (Fragment) OS=Aedes aegypti GN=AAEL003812 PE=4 SV=1
CCG026441.1	2041	1894	5709	39.8749032	97.78383248	1.294114935	Up	0	0	AAEL011180-PA OS=Aedes aegypti GN=AAEL011180 PE=4 SV=1
CCG019516.1	2313	1733	5737	32.1947904	86.7080129	1.429338067	Up	1.05E-12	2.59E-12	AAEL008854-PA OS=Aedes aegypti GN=AAEL008854 PE=4 SV=1

CCG013789.1	2433	402	5738	7.09980818	82.44578386	3.537593765	Up	0	0	AAEL009084-PA OS=Aedes aegypti GN=AAEL009084 PE=4 SV=1
CCG016307.1	5076	2173	5739	18.395042	39.52434011	1.103424404	Up	0	0	AAEL013896-PA (Fragment) OS=Aedes aegypti GN=AAEL013896 PE=4 SV=1
CCG000546.1	2427	1559	5740	27.601902	82.67841313	1.582743013	Up	6.25E-12	1.41E-11	AAEL001715-PA OS=Aedes aegypti GN=AAEL001715 PE=3 SV=1
CCG019832.1	451	13316	5757	1268.70285	446.2412403	-1.507458454	Down	0	0	AAEL014937-PA OS=Aedes aegypti GN=AAEL014937 PE=4 SV=1
CCG013129.1	3159	13824	5766	188.038497	63.80798476	-1.559219177	Down	0	0	AAEL000644-PA OS=Aedes aegypti GN=AAEL006446 PE=4 SV=1
CCG001350.1	3714	384	5772	4.44275124	54.32934128	3.612206288	Up	1.03E-12	2.55E-12	AAEL011287-PA OS=Aedes aegypti GN=AAEL011287 PE=4 SV=1
CCG013093.1	4650	1146	5794	10.5899604	43.55876465	2.040265844	Up	4.08E-12	9.43E-12	AAEL003733-PA OS=Aedes aegypti GN=AAEL003733 PE=4 SV=1
CCG027748.1	1483	660	5810	19.1234153	136.9572408	2.840313438	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_10459 PE=3 SV=1
CCG019953.1	2883	42723	5810	636.765859	70.45008257	-3.176089684	Down	0	0	Pyruvate kinase OS=Aedes aegypti GN=AAEL012576 PE=3 SV=1
CCG011270.1	5793	762	5814	5.65215567	35.08500279	2.633981372	Up	5.84E-13	1.50E-12	AAEL000832-PA OS=Aedes aegypti GN=AAEL000832 PE=4 SV=1
CCG028285.1	7677	1369	5817	7.66257219	26.48851061	1.789466061	Up	3.82E-12	8.84E-12	AAEL009959-PA OS=Aedes aegypti GN=AAEL009959 PE=4 SV=1
CCG013802.1	7643	1452	5817	8.16329382	26.60634515	1.704547055	Up	3.53E-12	8.22E-12	AAEL009959-PA OS=Aedes aegypti GN=AAEL009959 PE=4 SV=1
CCG008283.1	5661	1382	5818	10.49005	35.9277962	1.776078885	Up	5.79E-12	1.31E-11	AAEL000815-PA OS=Aedes aegypti GN=AAEL000815 PE=4 SV=1
CCG026148.1	1653	564	5819	14.6611799	123.0624395	3.069317382	Up	9.30E-13	2.31E-12	NMD protein OS=Aedes aegypti PE=2 SV=1
CCG018663.1	3673	29412	5828	344.085444	55.4687822	-2.633018904	Down	0	0	AAEL005422-PA OS=Aedes aegypti GN=AAEL005422 PE=3 SV=1
CCG028519.2	7646	874	5829	4.91179022	26.65077103	2.439856421	Up	0	0	AGAP001710-PA OS=Anopheles gambiae GN=AgaP_AGAP001710 PE=4 SV=2

CCG009106.1	22569	11800	5829	22.4663418	9.028835806	-1.315153349	Down	0	0	AAEL014026-PA OS=Aedes aegypti GN=AAEL014026 PE=4 SV=1
CCG009105.1	48632	1909	5830	1.6867335	4.190795229	1.312991985	Up	0	0	AAEL014030-PA OS=Aedes aegypti GN=AAEL014030 PE=4 SV=1
CCG024911.1	8430	1275	5851	6.49898121	24.26344928	1.900499175	Up	4.14E-12	9.55E-12	AAEL011479-PA OS=Aedes aegypti GN=AAEL011479 PE=4 SV=1
CCG015539.1	2088	2056	5854	42.3111946	98.01041773	1.211895685	Up	0	0	AAEL012821-PA OS=Aedes aegypti GN=AAEL012821 PE=4 SV=1
CCG012672.1	1819	43	5856	1.01577712	112.5429735	6.791748287	Up	8.46E-14	2.71E-13	AAEL001414-PB OS=Aedes aegypti GN=AAEL001414 PE=4 SV=1
CCG018861.1	4161	807	5856	8.33371205	49.1986707	2.561588179	Up	9.50E-13	2.36E-12	AAEL011879-PA OS=Aedes aegypti GN=AAEL011879 PE=4 SV=1
CCG007733.2	8303	265	5870	1.37142956	24.7145712	4.171609442	Up	4.94E-13	1.28E-12	AAEL004799-PA OS=Aedes aegypti GN=AAEL004799 PE=4 SV=1
CCG025144.1	2796	72	5873	1.10651677	73.42988537	6.052270125	Up	0	0	AAEL004831-PA OS=Aedes aegypti GN=AAEL004831 PE=4 SV=1
CCG026609.1	5848	13478	5880	99.0331879	35.14956693	-1.494405167	Down	0	0	-
CCG013218.2	3855	1739	5881	19.3837532	53.3306422	1.460116762	Up	4.99E-12	1.13E-11	AAEL003768-PA OS=Aedes aegypti GN=AAEL003768 PE=4 SV=1
CCG012125.1	1620	82	5898	2.17501126	127.2740225	5.870771306	Up	0	0	AAEL002554-PB OS=Aedes aegypti GN=AAEL002554 PE=4 SV=1
CCG007786.1	3132	2405	5930	32.9955976	66.18856362	1.004308415	Up	0	0	AAEL014721-PA OS=Aedes aegypti GN=AAEL014721 PE=4 SV=1
CCG002982.1	1951	1283	5933	28.2573908	106.3082809	1.911553816	Up	9.34E-13	2.32E-12	T-complex protein 1 subunit delta OS=Aedes aegypti GN=AAEL007702 PE=3 SV=1
CCG001971.1	13387	615	5939	1.97403352	15.50886724	2.973874921	Up	0	0	AAEL007813-PA OS=Aedes aegypti GN=AAEL007813 PE=4 SV=1
CCG002537.1	2897	266	5940	3.94544336	71.6783445	4.183277984	Up	4.09E-14	1.40E-13	AAEL001939-PA OS=Aedes aegypti GN=AAEL001939 PE=4 SV=1
CCG000265.1	11685	1463	5945	5.37995051	17.78578994	1.725060244	Up	0	0	Transcription-associated protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ013348 PE=4 SV=1
CCG008116.1	14590	973	5949	2.86563069	14.25406364	2.314448672	Up	3.18E-12	7.44E-12	AAEL001861-PA OS=Aedes aegypti GN=AAEL001861 PE=4 SV=1

CCG020441.1	1626	1070	5949	28.2765166	127.9008539	2.177349585	Up	1.81E-12	4.33E-12	AAEL002178-PA (Fragment) OS=Aedes aegypti GN=AAEL002178 PE=4 SV=1
CCG014886.1	17494	1246	5963	3.06049442	11.91586855	1.961047472	Up	2.10E-12	5.00E-12	AAEL013172-PA OS=Aedes aegypti GN=AAEL013172 PE=4 SV=1
CCG018108.1	3459	193	5986	2.39755964	60.4973243	4.65723273	Up	6.08E-14	2.02E-13	AAEL009475-PA (Fragment) OS=Aedes aegypti GN=AAEL009475 PE=3 SV=1
CCG007789.1	3078	1949	6010	27.2085811	68.25836368	1.326944104	Up	0	0	AAEL002528-PA OS=Aedes aegypti GN=AAEL002528 PE=4 SV=1
CCG011186.1	1919	1440	6012	32.2440948	109.5201459	1.764089401	Up	1.98E-12	4.73E-12	AAEL006946-PA OS=Aedes aegypti GN=AAEL006946 PE=3 SV=1
CCG021758.1	27156	2410	6012	3.81341363	7.739326851	1.021125067	Up	0	0	AAEL002272-PA OS=Aedes aegypti GN=AAEL002272 PE=4 SV=1
CCG015354.1	4201	1982	6015	20.2727955	50.05332892	1.303920979	Up	0	0	AAEL014597-PA OS=Aedes aegypti GN=AAEL014597 PE=4 SV=1
CCG027585.1	8925	2102	6031	10.1201549	23.62278624	1.22294777	Up	0	0	AAEL014637-PA OS=Aedes aegypti GN=AAEL014637 PE=4 SV=1
CCG020006.2	5399	638	6034	5.07773489	39.06987258	2.943799571	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_15518 PE=4 SV=1
CCG004244.1	4007	25655	6045	275.115683	52.73840357	-2.383112577	Down	0	0	AAEL006721-PA OS=Aedes aegypti GN=AAEL006721 PE=4 SV=1
CCG014463.1	936	11959	6058	549.011812	226.2577358	-1.278870071	Down	0	0	Activating transcription factor OS=Aedes aegypti PE=2 SV=1
CCG003010.1	4479	1243	6065	11.9248449	47.33689409	1.988994553	Up	6.46E-12	1.45E-11	AAEL000761-PA OS=Aedes aegypti GN=AAEL000761 PE=4 SV=1
CCG000981.1	2034	1308	6079	27.6324547	104.4795302	1.918784682	Up	4.37E-12	1.01E-11	AAEL008849-PA OS=Aedes aegypti GN=AAEL008849 PE=4 SV=1
CCG011398.1	43376	13241	6091	13.116983	4.908955732	-1.41794787	Down	0	0	AAEL005238-PA OS=Aedes aegypti GN=AAEL005238 PE=4 SV=1
CCG012136.1	924	2117	6093	98.4490565	230.5203251	1.227444675	Up	0	0	Glutathione transferase OS=Culex quinquefasciatus GN=CqGSTd1 PE=3 SV=1

CCG001125.1	1968	351	6094	7.66380937	108.2498672	3.820159768	Up	4.44E-15	1.60E-14	AAEL005744-PA OS=Aedes aegypti GN=AAEL005744 PE=4 SV=1
CCG004595.1	2039	18834	6097	396.906318	104.531934	-1.92485478	Down	0	0	Dihydrolipoyl dehydrogenase OS=Aedes aegypti GN=AAEL006928 PE=3 SV=1
CCG024716.2	3393	534	6103	6.76269917	62.87956473	3.216920147	Up	0	0	AAEL009381-PA OS=Aedes aegypti GN=AAEL009381 PE=4 SV=1
CCG028326.3	7197	668	6106	3.98829828	29.65891871	2.894620786	Up	9.28E-13	2.31E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007864 PE=4 SV=1
CCG006090.1	1429	6	6135	0.18041876	150.0832946	9.700198332	Up	8.38E-06	1.47E-05	AAEL005922-PA OS=Aedes aegypti GN=AAEL005922 PE=4 SV=1
CCG024982.1	1682	2342	6158	59.8306293	127.9863664	1.097033995	Up	0	0	Eukaryotic translation initiation factor 3 subunit E OS=Anopheles darlingi GN=AND_12720 PE=3 SV=1
CCG007440.1	879	442	6165	21.6070793	245.1851835	3.504295824	Up	0	0	AAEL006301-PA OS=Aedes aegypti GN=AAEL006301 PE=4 SV=1
CCG007802.1	1891	1553	6165	35.2892638	113.9702677	1.691356269	Up	3.26E-12	7.62E-12	AAEL005114-PA OS=Aedes aegypti GN=AAEL005114 PE=4 SV=1
CCG016287.1	3477	1161	6182	14.3479614	62.15475036	2.115018886	Up	3.30E-12	7.70E-12	AAEL000682-PA OS=Aedes aegypti GN=AAEL000682 PE=4 SV=1
CCG019403.1	7972	22660	6185	122.139261	27.12204489	-2.170989157	Down	0	0	PTANKL-interacting factor 1A OS=Culex quinquefasciatus GN=CpipJ_CPIJ011483 PE=4 SV=1
CCG014786.2	1407	1673	6201	51.093366	154.0698467	1.592376648	Up	3.18E-12	7.44E-12	Eukaryotic translation initiation factor OS=Forcipomyia taiwana PE=2 SV=1
CCG028615.1	2643	883	6214	14.3557607	82.1909693	2.517350102	Up	1.21E-13	3.73E-13	Phosphatase 2C gamma OS=Culex quinquefasciatus GN=CpipJ_CPIJ017206 PE=3 SV=1
CCG004275.1	4783	644	6234	5.78559672	45.56343245	2.977338765	Up	2.78E-13	7.74E-13	AAEL011960-PA OS=Aedes aegypti GN=AAEL011960 PE=4 SV=1
CCG001853.1	780	756	6235	41.6475889	279.4421227	2.746244624	Up	0	0	AAEL013533-PA OS=Aedes aegypti GN=AAEL013533 PE=4 SV=1

CCG000247.1	2108	1414	6235	28.8231522	103.3988879	1.842920644	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_22097 PE=3 SV=1
CCG015398.1	5751	679	6237	5.07328287	37.91249734	2.901681983	Up	0	0	AAEL004290-PA (Fragment) OS=Aedes aegypti GN=AAEL004290 PE=3 SV=1
CCG014631.1	4913	1571	6250	13.7401696	44.47165272	1.694486213	Up	3.96E-12	9.15E-12	AAEL004702-PA OS=Aedes aegypti GN=AAEL004702 PE=4 SV=1
CCG023214.2	7344	1302	6267	7.61800034	29.83163406	1.969358748	Up	2.82E-12	6.61E-12	AAEL000430-PA OS=Aedes aegypti GN=AAEL000430 PE=4 SV=1
CCG021436.1	4329	1457	6267	14.4622091	50.60834385	1.80708732	Up	5.09E-12	1.16E-11	AAEL004942-PA OS=Aedes aegypti GN=AAEL004942 PE=4 SV=1
CCG026419.1	1570	957	6276	26.1923796	139.7440414	2.415567726	Up	1.76E-12	4.21E-12	SZK active chromatin boundary protein OS=Culex quinquefasciatus GN=CqipJ_CPIJ008624 PE=4 SV=1
CCG003125.1	1288	66	6304	2.20186528	171.1001372	6.279970904	Up	0	0	AAEL007097-PA OS=Aedes aegypti GN=AAEL007097 PE=4 SV=1
CCG002171.1	943	149	6313	6.78949147	234.0313905	5.107254719	Up	2.33E-13	6.62E-13	-
CCG022362.1	1401	1088	6321	33.3697868	157.7239597	2.240787462	Up	2.82E-12	6.61E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_23192 PE=4 SV=1
CCG015607.1	3657	609	6377	7.15574744	60.95951079	3.090676952	Up	4.13E-13	1.09E-12	Diastolemin specific protein z50 OS=Culex quinquefasciatus GN=CqipJ_CPIJ004358 PE=4 SV=1
CCG022684.1	7513	18247	6378	104.361606	29.6770783	-1.814170083	Down	0	0	AMPK-gamma subunit OS=Aedes aegypti GN=AMPK-gamma PE=2 SV=1
CCG024834.3	5790	890	6384	6.60501966	38.54466993	2.544896614	Up	0	0	AAEL009357-PA (Fragment) OS=Aedes aegypti GN=AAEL009357 PE=4 SV=1
CCG013028.1	3241	1780	6394	23.5995457	68.96736244	1.547154707	Up	0	0	Histone deacetylase OS=Aedes aegypti GN=AAEL004586 PE=3 SV=1
CCG008775.1	1855	527	6395	12.2075742	120.5165391	3.303382696	Up	8.05E-13	2.01E-12	AAEL000739-PA OS=Aedes aegypti GN=AAEL000739 PE=4 SV=1

CCG025249.1	1809	1386	6397	32.9220852	123.6197327	1.90878143	Up	5.95E-12	1.34E-11	AAEL004697-PA OS=Aedes aegypti GN=AAEL004697 PE=4 SV=1
CCG000344.1	5028	1646	6397	14.0668622	44.47655062	1.660744351	Up	3.63E-12	8.44E-12	AAEL003671-PA OS=Aedes aegypti GN=AAEL003671 PE=4 SV=1
CCG006196.1	627	5	6402	0.34266136	356.9424049	10.02469207	Up	1.59E-05	2.76E-05	-
CCG027567.1	4581	575	6409	5.3934943	48.90801044	3.180778613	Up	1.93E-13	5.56E-13	AAEL000276-PA OS=Aedes aegypti GN=AAEL000276 PE=4 SV=1
CCG016034.1	2634	1461	6434	23.83401	85.39162975	1.841072962	Up	1.80E-12	4.32E-12	AAEL002083-PA OS=Aedes aegypti GN=AAEL002083 PE=3 SV=1
CCG015656.1	3543	1230	6436	14.9175201	63.50309605	2.089819214	Up	3.75E-12	8.70E-12	Neuronal protein a OS=Culex quinquefasciatus GN=CpipJ_CPIJ013227 PE=4 SV=1
CCG027262.2	6575	1059	6451	6.92090478	34.2989876	2.309133431	Up	4.65E-12	1.06E-11	AAEL001322-PA OS=Aedes aegypti GN=AAEL001322 PE=4 SV=1
CCG025556.1	3384	366	6466	4.64743584	66.79675461	3.845271158	Up	0	0	AAEL011023-PA OS=Aedes aegypti GN=AAEL011023 PE=4 SV=1
CCG017111.1	2293	19217	6484	360.117484	98.85279834	-1.865113932	Down	0	0	AAEL000683-PA OS=Aedes aegypti GN=AAEL006834 PE=3 SV=1
CCG003437.1	459	2470	6491	231.231469	494.3663933	1.096242918	Up	0	0	AAEL012585-PA OS=Aedes aegypti GN=AAEL012585 PE=2 SV=1
CCG021496.1	3909	1322	6496	14.5321026	58.09387718	1.999142659	Up	0	0	AAEL010234-PA (Fragment) OS=Aedes aegypti GN=AAEL010234 PE=4 SV=1
CCG018436.1	12378	2184	6497	7.58166912	18.34900018	1.275114053	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_00689 PE=4 SV=1
CCG007316.1	4515	1330	6503	12.6577513	50.35075833	1.991992382	Up	4.68E-12	1.07E-11	AAEL009621-PA OS=Aedes aegypti GN=AAEL009621 PE=4 SV=1
CCG012673.1	1506	158	6507	4.50811293	151.0448253	5.066309294	Up	0	0	AAEL001420-PA OS=Aedes aegypti GN=AAEL001420 PE=4 SV=1
CCG007983.1	4521	2408	6509	22.8867775	50.33033035	1.136913726	Up	0	0	AAEL004984-PA OS=Aedes aegypti GN=AAEL004984 PE=4 SV=1
CCG005088.1	2721	711	6522	11.2280343	83.79194455	2.89970618	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008798 PE=4 SV=1

CCG028463.2	8107	1835	6522	9.72609634	28.12358223	1.531847582	Up	0	0	AAEL009073-PA OS=Reedes aegypti GN=AAEL009073 PE=4 SV=1
CCG018564.1	1813	2172	6543	51.4783583	126.1621649	1.293241371	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_11769 PE=4 SV=1
CCG008612.2	2542	607	6554	10.2606723	90.13239416	3.134920455	Up	0	0	AAEL010317-PA OS=Reedes aegypti GN=AAEL010317 PE=3 SV=1
CCG002938.1	38569	2534	6568	2.82313017	5.953121985	1.076350806	Up	0	0	AAEL003733-PA OS=Reedes aegypti GN=AAEL003733 PE=4 SV=1
CCG026319.1	13659	145093	6570	456.446863	16.81498487	-4.762627534	Down	0	0	AAEL002759-PB OS=Reedes aegypti GN=AAEL002759 PE=3 SV=1
CCG017011.1	29345	1263	6571	1.84940449	7.827937866	2.081571507	Up	2.88E-12	6.75E-12	-
CCG014174.1	2556	1565	6578	26.3097162	89.96695798	1.773799555	Up	1.11E-12	2.73E-12	AAEL002834-PA OS=Reedes aegypti GN=AAEL002834 PE=4 SV=1
CCG027923.1	3450	967	6600	12.0439807	66.87670339	2.473191433	Up	2.56E-13	7.18E-13	AGAP003537-PB OS=Anopheles gambiae GN=AgaP_AGAP003537 PE=4 SV=1
CCG016224.1	3954	759	6619	8.24836332	58.52019068	2.826754684	Up	1.54E-12	3.70E-12	AAEL006205-PB OS=Reedes aegypti GN=AAEL006205 PE=4 SV=1
CCG020172.1	1737	4	6623	0.09895161	133.2922666	10.39558235	Up	2.95E-05	5.04E-05	AAEL003404-PA OS=Reedes aegypti GN=AAEL003404 PE=4 SV=1
CCG018446.1	2481	16338	6626	282.966354	93.36297537	-1.599708076	Down	0	0	AAEL004739-PA OS=Reedes aegypti GN=AAEL004739 PE=3 SV=1
CCG013468.1	1695	1130	6646	28.6464897	137.0694439	2.258476724	Up	2.84E-12	6.65E-12	AAEL005528-PA OS=Reedes aegypti GN=AAEL005528 PE=4 SV=1
CCG008277.1	1089	71918	6673	2837.73864	214.2117364	-3.727632288	Down	0	0	AAEL002417-PC OS=Reedes aegypti GN=AAEL002417 PE=4 SV=1
CCG004090.1	738	1392	6675	81.0486051	316.1876659	1.963921829	Up	2.91E-12	6.82E-12	Nucleolar protein NOP5 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014566 PE=4 SV=1
CCG006552.1	2811	2231	6686	34.1036919	83.14871523	1.285766045	Up	0	0	AAEL000994-PA OS=Reedes aegypti GN=AAEL000994 PE=4 SV=1
CCG007151.1	1490	2229	6694	64.2815694	157.0541642	1.288785138	Up	0	0	AAEL004484-PA OS=Reedes aegypti GN=AAEL004484 PE=4 SV=1

CCG013416.1	16232	1400	6698	3.70611314	14.4252426	1.960616753	Up	8.19E-13	2.05E-12	AAEL010899-PA OS=Aedes aegypti GN=AAEL010899 PE=4 SV=1
CCG022863.1	810	536	6734	28.4342935	290.6284392	3.353472019	Up	0	0	AAEL012865-PA OS=Aedes aegypti GN=AAEL012865 PE=4 SV=1
CCG022785.1	7515	557	6735	3.18484926	31.3298728	3.298241917	Up	5.29E-13	1.37E-12	AAEL000295-PA OS=Aedes aegypti GN=AAEL000295 PE=4 SV=1
CCG015560.1	2324	987	6735	18.2491945	101.3098081	2.47286916	Up	3.78E-12	8.77E-12	AAEL005567-PA OS=Aedes aegypti GN=AAEL005567 PE=2 SV=1
CCG008966.1	16556	12482	6746	32.3960031	14.24429422	-1.18543169	Down	0	0	TXR DNA-binding protein 45 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014288 PE=4 SV=1
CCG009398.1	4642	1116	6754	10.3305092	50.86346431	2.299718356	Up	3.58E-12	8.33E-12	AAEL006320-PA OS=Aedes aegypti GN=AAEL006320 PE=4 SV=1
CCG013600.3	3330	16056	6759	207.183801	70.95585366	-1.545917599	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_21730 PE=3 SV=1
CCG017035.1	3947	1464	6770	15.9381027	59.96137161	1.911553484	Up	4.30E-12	9.90E-12	AAEL006205-PA OS=Aedes aegypti GN=AAEL006205 PE=4 SV=1
CCG011048.1	9092	16922	6772	79.9751264	26.03799497	-1.618933008	Down	0	0	AAEL011604-PA OS=Aedes aegypti GN=AAEL011604 PE=4 SV=1
CCG021500.1	753	4	6784	0.22825888	314.9494683	10.43023366	Up	2.95E-05	5.04E-05	SV kDa salivary gland antigen 30k-3 OS=Aedes albopictus PE=2 SV=1
CCG028513.1	45562	1197	6784	1.12889628	5.205147922	2.205026221	Up	7.99E-13	2.00E-12	Putative methoprene-tolerant mutant OS=Aedes aegypti PE=2 SV=1
CCG003634.1	3302	2052	6815	26.7031785	72.15041072	1.43399613	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_11093 PE=4 SV=1
CCG023382.1	3741	705	6818	8.09774469	63.71171639	2.975966641	Up	7.66E-13	1.92E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009921 PE=4 SV=1
CCG027773.1	3045	789	6821	11.1340298	78.30883608	2.814199261	Up	0	0	AAEL001032-PA OS=Aedes aegypti GN=AAEL001032 PE=4 SV=1
CCG016658.1	4899	813	6826	7.1309235	48.70896045	2.772026361	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001658 PE=4 SV=1

CCG023105.1	18387	2187	6833	5.1109376	12.99123866	1.34587911	Up	0	0	AAEL008015-PA OS=Aedes aegypti GN=AAEL008015 PE=4 SV=1
CCG027735.1	2975	1447	6843	20.8999012	80.40991192	1.943877229	Up	6.55E-12	1.47E-11	AAEL009895-PA (Fragment) OS=Aedes aegypti GN=AAEL009895 PE=4 SV=1
CCG008582.1	16927	677	6858	1.7185863	14.16339943	3.042873369	Up	4.03E-13	1.07E-12	AAEL007698-PA OS=Aedes aegypti GN=PIWI4 PE=4 SV=1
CCG024974.1	3195	1782	6883	23.9662182	75.31074148	1.651853377	Up	0	0	AAEL002705-PA OS=Aedes aegypti GN=AAEL002705 PE=4 SV=1
CCG025523.1	9764	2183	6894	9.60701871	24.68274888	1.361342372	Up	0	0	AAEL008579-PA OS=Aedes aegypti GN=AAEL008579 PE=4 SV=1
CCG014706.2	11574	1137	6895	4.22123624	20.82575759	2.302631501	Up	4.86E-12	1.11E-11	PolyQ domain-containing hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG007906.1	1686	2477	6901	63.1293195	143.0884152	1.180524768	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009698 PE=4 SV=1
CCG006070.1	2229	2621	6901	50.5265475	108.2310758	1.099001283	Up	0	0	AAEL007042-PA OS=Aedes aegypti GN=AAEL007042 PE=4 SV=1
CCG007533.1	2068	11275	6903	234.276479	116.6909983	-1.00551884	Down	0	0	Ribose-phosphate pyrophosphokinase OS=Aedes aegypti GN=AAEL004178 PE=3 SV=1
CCG021341.1	4974	971	6909	8.38834184	48.55784765	2.533246915	Up	4.63E-12	1.06E-11	AAEL000181-PA OS=Aedes aegypti GN=AAEL000181 PE=4 SV=1
CCG023237.1	2388	1858	6909	33.4329007	101.1418485	1.597039614	Up	0	0	AAEL012827-PA OS=Aedes aegypti GN=AAEL012827 PE=3 SV=1
CCG026844.1	3440	1466	6912	18.312102	70.24174682	1.939531318	Up	1.74E-12	4.17E-12	AAEL005528-PA OS=Aedes aegypti GN=AAEL005528 PE=4 SV=1
CCG003448.1	1879	2549	6917	58.2915665	128.6888773	1.142528287	Up	0	0	AAEL005877-PA OS=Aedes aegypti GN=AAEL005877 PE=4 SV=1
CCG002972.1	5871	2497	6933	18.2754943	41.28184855	1.175597143	Up	0	0	AAEL009706-PA OS=Aedes aegypti GN=AAEL009706 PE=4 SV=1

CCG010771.1	4245	1267	6936	12.8251246	57.11910664	2.155000578	Up	3.44E-12	8.01E-12	AAEL007538-PA (Fragment) OS=Aedes aegypti GN=AAEL007538 PE=4 SV=1
CCG018659.2	37887	2507	6942	2.84332686	6.405372749	1.171702573	Up	0	0	AAEL001869-PA OS=Aedes aegypti GN=AAEL001869 PE=4 SV=1
CCG025447.1	1737	807	6947	19.9634864	139.8129814	2.808062723	Up	9.50E-13	2.36E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ004712 PE=4 SV=1
CCG005554.1	6110	2006	6950	14.1075757	39.76432464	1.495004576	Up	0	0	AAEL003275-PA OS=Aedes aegypti GN=AAEL003275 PE=4 SV=2
CCG024227.1	7318	1582	6966	9.28916646	33.2767636	1.840894077	Up	1.12E-12	2.76E-12	AAEL001910-PA OS=Aedes aegypti GN=AAEL001910 PE=4 SV=1
CCG009741.1	3399	1073	6969	13.5647323	71.6752665	2.401614783	Up	3.38E-12	7.88E-12	AAEL014212-PA OS=Aedes aegypti GN=AAEL014212 PE=4 SV=1
CCG023537.1	2835	1873	6976	28.3888229	86.02078969	1.599362344	Up	0	0	AAEL013334-PA OS=Aedes aegypti GN=AAEL013334 PE=4 SV=1
CCG016238.1	5792	2176	6977	16.1433257	42.11047946	1.383241481	Up	0	0	AAEL004327-PA OS=Aedes aegypti GN=AAEL004327 PE=3 SV=1
CCG020044.1	1431	1274	7003	38.2553752	171.0781357	2.160921014	Up	1.62E-12	3.89E-12	AAEL013653-PA OS=Aedes aegypti GN=AAEL013653 PE=4 SV=1
CCG025920.2	4296	827	7008	8.27187396	57.0269096	2.785356744	Up	1.96E-12	4.67E-12	AAEL012290-PA OS=Aedes aegypti GN=AAEL012290 PE=4 SV=1
CCG001115.1	2247	1294	7017	24.7453656	109.1687709	2.141329949	Up	3.05E-12	7.15E-12	AAEL000857-PA OS=Aedes aegypti GN=AAEL000857 PE=4 SV=1
CCG012679.1	4407	1358	7022	13.2409575	55.7016155	2.07271172	Up	5.15E-12	1.17E-11	AAEL007258-PA OS=Aedes aegypti GN=AAEL007258 PE=4 SV=1
CCG013370.1	4191	571	7027	5.85438284	58.6141281	3.323659452	Up	0	0	AAEL006249-PA OS=Aedes aegypti GN=AAEL006249 PE=4 SV=1
CCG026522.1	4248	1271	7036	12.8565284	57.90170324	2.171104656	Up	5.38E-12	1.22E-11	AAEL009934-PA OS=Aedes aegypti GN=AAEL009934 PE=4 SV=1
CCG004459.2	41423	1677	7040	1.73961917	5.941295137	1.772005944	Up	3.09E-12	7.24E-12	AAEL009357-PA (Fragment) OS=Aedes aegypti GN=AAEL009357 PE=4 SV=1
CCG011648.1	5901	1514	7087	11.0246023	41.98429207	1.929123048	Up	3.22E-12	7.53E-12	AAEL004839-PA OS=Aedes aegypti GN=AAEL004839 PE=3 SV=1

CCG012664.1	3327	1034	7092	13.3545854	74.51881541	2.480269554	Up	3.89E-12	8.99E-12	Uncharacterized protein OS=Anopheles darlingi GN=AND_06456 PE=3 SV=1
CCG004387.1	3982	1472	7094	15.8843419	62.27875827	1.971134863	Up	4.36E-12	1.00E-11	AAEL010698-PA (Fragment) OS=Aedes aegypti GN=AAEL010698 PE=4 SV=1
CCG021737.1	834	2354	7097	121.283879	297.4806838	1.29440819	Up	0	0	60S ribosomal protein L13a (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG016817.1	4426	204	7103	1.98053002	56.10226839	4.824100632	Up	6.59E-14	2.17E-13	AAEL014694-PA OS=Aedes aegypti GN=AAEL014694 PE=4 SV=1
CCG026044.1	2661	31612	7118	510.469467	93.51109134	-2.448615272	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009078 PE=4 SV=1
CCG007358.1	5300	2152	7125	17.4473338	46.9957966	1.42952514	Up	0	0	AAEL000564-PA OS=Aedes aegypti GN=AAEL000564 PE=4 SV=1
CCG003528.1	3663	361	7131	4.23480049	68.05554782	4.006346866	Up	1.35E-13	4.08E-13	AAEL000388-PA OS=Aedes aegypti GN=AAEL000388 PE=3 SV=1
CCG023895.1	1270	1302	7139	44.0524366	196.5095574	2.157305757	Up	2.82E-12	6.61E-12	AAEL010668-PA OS=Aedes aegypti GN=AAEL010668 PE=4 SV=1
CCG023812.1	1945	905	7157	19.9936297	128.6356745	2.685678486	Up	0	0	AAEL012126-PA OS=Aedes aegypti GN=AAEL012126 PE=4 SV=1
CCG016471.1	7642	1908	7170	10.7283766	32.799116	1.612225151	Up	0	0	AAEL005231-PA OS=Aedes aegypti GN=AAEL005231 PE=4 SV=1
CCG012781.1	2112	654	7173	13.305969	118.7290338	3.157527294	Up	1.01E-12	2.50E-12	Adenosylhomocysteinase OS=Aedes aegypti GN=AAEL008341 PE=2 SV=1
CCG012782.1	3104	2430	7184	33.6393219	80.90858902	1.26614424	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_03468 PE=4 SV=1
CCG001704.1	3357	1431	7209	18.3168574	75.07125923	2.035088681	Up	5.80E-12	1.31E-11	AAEL001540-PA OS=Aedes aegypti GN=AAEL001540 PE=4 SV=1
CCG016920.1	3180	700	7230	9.45874661	79.4806104	3.070882024	Up	0	0	Dab2-interacting protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007375 PE=4 SV=1

CCG018110.1	4158	173	7251	1.78782205	60.96259377	5.091649235	Up	3.60E-13	9.69E-13	AAEL009490-PA (Fragment) OS=Aedes aegypti GN=AAEL009490 PE=3 SV=1
CCG016919.1	1227	2004	7272	70.1803978	207.1854839	1.561782895	Up	0	0	AAEL001005-PA OS=Aedes aegypti GN=AAEL001005 PE=3 SV=1
CCG021760.1	2860	1969	7290	29.5830096	89.10693625	1.590768907	Up	0	0	AAEL008374-PA OS=Aedes aegypti GN=AAEL008374 PE=4 SV=1
CCG023775.1	6679	1638	7305	10.5381682	38.2347974	1.85926212	Up	7.06E-12	1.58E-11	DNA-directed RNA polymerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ016666 PE=3 SV=1
CCG019398.1	516	1906	7332	158.721539	496.7327234	1.64597187	Up	0	0	60S ribosomal protein L21 OS=Aedes aegypti GN=AAEL007715 PE=2 SV=1
CCG016849.1	3587	1299	7339	15.5611055	71.52461478	2.20049527	Up	7.92E-13	1.99E-12	AAEL008862-PA OS=Aedes aegypti GN=AAEL008862 PE=3 SV=1
CCG021605.1	4380	2785	7345	27.3220801	58.622955	1.101398367	Up	0	0	AAEL008389-PA (Fragment) OS=Aedes aegypti GN=AAEL008389 PE=4 SV=1
CCG001100.1	2097	2376	7361	48.6867379	122.7123869	1.333680139	Up	0	0	AAEL013538-PA OS=Aedes aegypti GN=AAEL013538 PE=4 SV=1
CCG010807.5	16644	31762	7371	81.9998024	15.4817026	-2.405056292	Down	0	0	AAEL013559-PA OS=Aedes aegypti GN=AAEL013559 PE=4 SV=1
CCG004997.1	1164	497	7373	18.347043	221.4324524	3.593247204	Up	7.04E-14	2.29E-13	AAEL000604-PA OS=Aedes aegypti GN=AAEL000604 PE=4 SV=1
CCG011547.1	1395	70899	7373	2183.87901	184.7651431	-3.563128413	Down	0	0	Citrate synthase OS=Anopheles darlingi GN=AND_04269 PE=3 SV=1
CCG014217.1	5548	887	7380	6.86989088	46.50181734	2.75892801	Up	1.39E-12	3.35E-12	Zinc finger protein 650 OS=Culex quinquefasciatus GN=CpipJ_CPIJ009486 PE=4 SV=1
CCG022466.2	24426	2906	7381	5.11217755	10.5636224	1.047094791	Up	0	0	AAEL008881-PA OS=Aedes aegypti GN=AAEL008881 PE=4 SV=1
CCG018614.1	2908	39672	7395	586.208842	88.89836889	-2.721185874	Down	0	0	AAEL010608-PA (Fragment) OS=Aedes aegypti GN=AAEL010608 PE=4 SV=1

CCG008286.1	31410	2066	7414	2.82634421	8.251533397	1.545725065	Up	0	0	AAEL013131-PA OS=Aedes aegypti GN=AAEL013131 PE=4 SV=1
CCG001636.1	1143	499	7478	18.7593155	228.7121555	3.607853954	Up	0	0	AAEL001216-PA OS=Aedes aegypti GN=AAEL001216 PE=2 SV=1
CCG025748.1	3795	1811	7480	20.5054517	68.90327016	1.748564928	Up	0	0	AAEL012989-PA OS=Aedes aegypti GN=AAEL012989 PE=4 SV=1
CCG017415.3	2893	1712	7482	25.4283393	90.41058652	1.830054468	Up	5.08E-12	1.16E-11	AAEL011089-PA OS=Aedes aegypti GN=AAEL011089 PE=4 SV=1
CCG022723.1	5306	2236	7486	18.107864	49.32108178	1.445588064	Up	0	0	AAEL004838-PA OS=Aedes aegypti GN=AAEL004838 PE=4 SV=1
CCG009547.1	4117	1073	7502	11.1990588	63.70099401	2.507938391	Up	3.38E-12	7.87E-12	AAEL009189-PA OS=Aedes aegypti GN=AAEL009189 PE=4 SV=1
CCG006708.1	2628	298	7504	4.87251937	99.81998056	4.356588796	Up	3.92E-13	1.05E-12	AAEL008098-PA OS=Aedes aegypti GN=PIWI2 PE=4 SV=1
CCG013829.2	1800	223	7519	5.32347268	146.0284906	4.77773839	Up	0	0	Rap55 OS=Culex quinquefasciatus
CCG025916.1	15700	889	7524	2.43312701	16.75325315	2.783557729	Up	0	0	AAEL007823-PA OS=Aedes aegypti GN=AAEL007823 PE=4 SV=1
CCG003898.4	3426	2825	7549	35.4318448	77.02861394	1.120347883	Up	0	0	AAEL009437-PA (Fragment) OS=Aedes aegypti GN=AAEL009437 PE=4 SV=1
CCG025503.1	2493	756	7555	13.0305332	105.9405459	3.02328682	Up	0	0	AAEL012918-PA OS=Aedes aegypti GN=AAEL012918 PE=4 SV=1
CCG019279.1	3303	675	7563	8.78128092	80.04524591	3.188312415	Up	1.92E-12	4.59E-12	AAEL005453-PA (Fragment) OS=Aedes aegypti GN=AAEL005453 PE=4 SV=1
CCG005054.1	5137	2028	7563	16.9637185	51.46767515	1.60121417	Up	0	0	AAEL001454-PA OS=Aedes aegypti GN=AAEL001454 PE=4 SV=1
CCG020660.1	1701	18673	7564	471.707145	155.4523254	-1.601419257	Down	0	0	AAEL004034-PA OS=Aedes aegypti GN=CYP4G36 PE=3 SV=1
CCG004898.1	2395	1006	7570	18.0490827	110.494428	2.613976199	Up	4.55E-12	1.05E-11	AAEL001004-PA OS=Aedes aegypti GN=AAEL001004 PE=4 SV=1
CCG025305.1	23693	1891	7590	3.42952636	11.19880643	1.707263726	Up	0	0	AAEL009719-PA OS=Aedes aegypti GN=AAEL009719 PE=4 SV=1

CCG015033.1	1610	2734	7591	72.968481	164.825018	1.175589912	Up	0	0	AAEL007160-PA OS=Aedes aegypti GN=AAEL007160 PE=4 SV=1
CCG024840.1	32234	1612	7606	2.14888665	8.248825871	1.940599399	Up	1.56E-12	3.75E-12	Serine/threonine-protein kinase D OS=Aedes aegypti GN=AAEL002892 PE=3 SV=1
CCG020268.1	1251	1350	7616	46.3702172	212.8235299	2.198387275	Up	0	0	AAEL006353-PA OS=Aedes aegypti GN=AAEL008353 PE=2 SV=1
CCG026083.1	1398	941	7643	28.9231189	191.1202499	2.724185606	Up	0	0	AAEL006830-PA OS=Aedes aegypti GN=AAEL006830 PE=4 SV=1
CCG001101.1	11202	12574	7693	48.2325873	24.00767927	-1.006512251	Down	0	0	AAEL013530-PA OS=Aedes aegypti GN=AAEL013530 PE=3 SV=1
CCG007937.2	49617	2916	7702	2.52533902	5.426540252	1.103555607	Up	0	0	AAEL010184-PA OS=Aedes aegypti GN=AAEL010184 PE=4 SV=1
CCG023566.1	1568	55837	7717	1530.1665	172.0491211	-3.152796216	Down	0	0	-
CCG025941.1	4046	895	7730	9.50516868	66.78879868	2.81282203	Up	9.98E-13	2.47E-12	AAEL011876-PA OS=Aedes aegypti GN=AAEL011876 PE=4 SV=1
CCG025019.1	15173	2502	7731	7.08563079	17.81206338	1.329886452	Up	0	0	AAEL014098-PA OS=Aedes aegypti GN=AAEL014098 PE=4 SV=1
CCG019436.1	2793	2120	7737	32.6157671	96.83930805	1.570023213	Up	0	0	AAEL003800-PA OS=Aedes aegypti GN=AAEL003800 PE=4 SV=1
CCG013430.1	276	22	7743	3.42512377	980.731656	8.161558512	Up	8.38E-11	1.80E-10	Ribosomal protein L37 OS=Aedes aegypti PE=3 SV=1
CCG026722.1	1443	61476	7749	1830.63576	187.7281266	-3.285628045	Down	0	0	AAEL001673-PA OS=Aedes aegypti GN=AAEL001673 PE=3 SV=1
CCG012529.1	1839	26471	7779	618.516501	147.874081	-2.064442881	Down	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008687 PE=4 SV=1
CCG023268.1	8013	706	7786	3.78592695	33.96794496	3.16545546	Up	0	0	AAEL006187-PA OS=Aedes aegypti GN=AAEL006187 PE=4 SV=1
CCG018341.1	3620	1424	7787	16.9030116	75.19892299	2.153431684	Up	4.81E-12	1.10E-11	AAEL013173-PA OS=Aedes aegypti GN=AAEL013173 PE=4 SV=1
CCG025165.1	1305	2198	7787	72.3735453	208.5977787	1.527189444	Up	0	0	Lipid storage droplets surface-binding protein 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003879 PE=4 SV=1

CCG016031.1	3900	2839	7811	31.2797632	70.01515381	1.162437615	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015901 PE=4 SV=1
CCG002156.2	3698	2732	7813	31.7450825	73.85857664	1.218232334	Up	0	0	AAEL001919-PA OS=Aedes aegypti GN=AAEL001919 PE=4 SV=1
CCG021633.1	6440	1931	7861	12.8842481	42.67189654	1.727677889	Up	0	0	AGAP012219-PA OS=Anopheles gambiae GN=AGAP012219 PE=4 SV=4
CCG024675.1	1860	14030	7883	324.12117	148.1591913	-1.129385125	Down	0	0	AAEL015076-PA OS=Aedes aegypti GN=AAEL015076 PE=4 SV=1
CCG019309.1	1548	0	7889	0.001	178.1562309	17.44278342	Up	4.44E-16	1.65E-15	AAEL013703-PA (Fragment) OS=Aedes aegypti GN=AAEL013703 PE=3 SV=1
CCG009481.1	14959	2758	7889	7.92235631	18.43611508	1.218533185	Up	0	0	AAEL004981-PA OS=Aedes aegypti GN=AAEL004981 PE=4 SV=1
CCG013152.1	5808	1543	7893	11.4156853	47.50786477	2.057148891	Up	1.47E-12	3.54E-12	AAEL013131-PA OS=Aedes aegypti GN=AAEL013131 PE=4 SV=1
CCG021104.1	1809	2807	7896	66.6755362	152.5873706	1.19440613	Up	0	0	AAEL001959-PA OS=Aedes aegypti GN=AAEL001959 PE=4 SV=1
CCG004911.1	2523	2097	7901	35.7144405	109.4749682	1.616021604	Up	0	0	AGAP010331-PA OS=Anopheles gambiae GN=AGAP010331 PE=3 SV=3
CCG007993.1	4200	146	7903	1.49370982	65.77982413	5.460673338	Up	9.33E-15	3.34E-14	AAEL011312-PA OS=Aedes aegypti GN=AAEL011312 PE=4 SV=1
CCG009980.1	16735	1766	7907	4.5344817	16.51718521	1.864958285	Up	1.35E-12	3.28E-12	AAEL001165-PA OS=Aedes aegypti GN=AAEL001165 PE=4 SV=1
CCG002534.1	1794	2523	7924	60.4306803	154.4087989	1.353401878	Up	0	0	AAEL001052-PA OS=Aedes aegypti GN=AAEL001052 PE=3 SV=1
CCG004231.1	2970	2684	7932	38.8319083	93.36331696	1.26561321	Up	0	0	AAEL004886-PA OS=Aedes aegypti GN=AAEL004886 PE=4 SV=1
CCG005650.1	2030	26502	7961	560.977294	137.0949958	-2.032766467	Down	0	0	AAEL008073-PA OS=Aedes aegypti GN=AAEL008073 PE=4 SV=1
CCG022749.1	3360	2592	7987	33.148081	83.09873708	1.325901199	Up	0	0	AAEL002646-PA OS=Aedes aegypti GN=AAEL002646 PE=4 SV=1

CCG004134.1	3586	2471	7992	29.6090949	77.91035916	1.395774776	Up	0	0	AAEL003466-PA OS=Aedes aegypti GN=AAEL003466 PE=3 SV=1
CCG015276.2	30756	19800	7997	27.6629193	9.089652079	-1.605656433	Down	0	0	AAEL009872-PA OS=Aedes aegypti GN=AAEL009872 PE=4 SV=1
CCG025355.3	15242	2023	7998	5.70317367	18.34380643	1.685456165	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_14521 PE=4 SV=1
CCG008930.1	3144	1265	8010	17.2890313	89.06354865	2.364978062	Up	6.04E-12	1.36E-11	AAEL011240-PA OS=Aedes aegypti GN=AAEL011240 PE=4 SV=1
CCG010790.1	1920	6	8053	0.13428042	146.6244806	10.09266132	Up	8.38E-06	1.47E-05	AAEL002652-PA OS=Aedes aegypti GN=AAEL002652 PE=4 SV=1
CCG008616.1	3073	1560	8062	21.8134676	91.71286278	2.071904958	Up	7.56E-13	1.90E-12	AAEL010311-PA OS=Aedes aegypti GN=AAEL010311 PE=4 SV=1
CCG017078.1	2111	192	8067	3.90818998	133.5899662	5.095167244	Up	0	0	Importin subunit alpha OS=Aedes aegypti GN=AAEL012960 PE=3 SV=1
CCG025991.2	2582	1683	8080	28.0085451	109.3969312	1.96563332	Up	1.60E-12	3.86E-12	AAEL010826-PA OS=Aedes aegypti GN=AAEL010826 PE=3 SV=1
CCG018403.1	3519	1492	8082	18.2184837	80.28780701	2.13977802	Up	2.02E-12	4.82E-12	AAEL004500-PA OS=Aedes aegypti GN=AAEL004500 PE=4 SV=1
CCG012666.1	1239	2055	8086	71.2694145	228.1457837	1.678601014	Up	0	0	Carboxypeptidase B OS=Aedes polynesiensis GN=CPB-I PE=3 SV=1
CCG009878.1	3765	1444	8103	16.4802913	75.23689686	2.190698602	Up	3.35E-12	7.81E-12	AAEL000388-PA OS=Aedes aegypti GN=AAEL000388 PE=3 SV=1
CCG009226.1	21237	2669	8106	5.40030238	13.34330609	1.305004074	Up	0	0	AAEL002987-PA (Fragment) OS=Aedes aegypti GN=AAEL002987 PE=3 SV=1
CCG025176.1	2985	2051	8121	29.5245982	95.10759319	1.6876431	Up	0	0	AAEL000624-PA OS=Aedes aegypti GN=AAEL000624 PE=4 SV=1
CCG009514.1	63359	3244	8123	2.20006343	4.48185865	1.026552029	Up	0	0	AAEL001389-PA OS=Aedes aegypti GN=AAEL001389 PE=4 SV=1

CCG018319.2	1301	31259	8137	1032.42962	218.6437341	-2.239389526	Down	0	0	Cytochrome b-c1 complex subunit Rieske, mitochondrial OS=Aedes aegypti GN=AAEL003675 PE=3 SV=1
CCG018875.1	2616	804	8139	13.2062946	108.7635377	3.041897346	Up	2.72E-13	7.60E-13	AAEL001791-PA (Fragment) OS=Aedes aegypti GN=AAEL001791 PE=4 SV=1
CCG000847.1	3134	1845	8141	25.2964775	90.80897613	1.843898406	Up	0	0	AAEL005277-PA OS=Aedes aegypti GN=AAEL005277 PE=4 SV=1
CCG028554.1	2430	1555	8144	27.4970935	117.1605786	2.091136185	Up	2.51E-12	5.91E-12	AAEL011282-PA OS=Aedes aegypti GN=AAEL011282 PE=4 SV=1
CCG013965.1	5382	2308	8144	18.4270062	52.89858901	1.521407541	Up	0	0	AAEL003955-PA OS=Aedes aegypti GN=AAEL003955 PE=4 SV=1
CCG021097.1	297	531	8147	76.824677	958.9396662	3.641798346	Up	2.40E-13	6.77E-13	AAEL001511-PA OS=Aedes aegypti GN=AAEL001511 PE=4 SV=1
CCG013466.3	969	1889	8148	83.7665931	293.9525688	1.811136482	Up	0	0	AAEL005515-PE OS=Aedes aegypti GN=AAEL005515 PE=4 SV=1
CCG010191.3	3270	2471	8154	32.4704019	87.17118924	1.42472615	Up	0	0	AAEL001218-PA OS=Aedes aegypti GN=AAEL001218 PE=3 SV=1
CCG024106.1	2384	907	8220	16.3479653	120.5356691	2.882277142	Up	0	0	AAEL000193-PA OS=Aedes aegypti GN=AAEL000193 PE=4 SV=1
CCG027343.1	3846	1945	8273	21.730664	75.19756207	1.790953631	Up	0	0	AAEL000293-PA OS=Aedes aegypti GN=AAEL000293 PE=4 SV=1
CCG018664.1	36469	2761	8283	3.2531585	7.939877883	1.287275705	Up	0	0	AAEL002037-PA OS=Aedes aegypti GN=AAEL002037 PE=4 SV=1
CCG017097.1	7377	26021	8377	151.567773	39.69709699	-1.932857624	Down	0	0	AAEL006307-PA OS=Aedes aegypti GN=AAEL006307 PE=4 SV=1
CCG004782.1	3351	459	8419	5.88573804	87.82862792	3.899396027	Up	5.70E-13	1.47E-12	-
CCG009685.1	2371	160470	8482	2908.20469	125.0595123	-4.539442113	Down	0	0	AAEL001593-PC OS=Aedes aegypti GN=AAEL001593 PE=3 SV=1
CCG027416.1	3669	15409	8519	180.463516	81.16913596	-1.152704039	Down	0	0	Phosphorylase OS=Aedes aegypti GN=AAEL000703 PE=3 SV=1
CCG003710.1	351	1995	8569	244.229688	853.4400959	1.805051309	Up	0	0	60S ribosomal protein L30 OS=Aedes aegypti GN=AAEL005085 PE=3 SV=1

CCG022453.1	10289	24686	8581	103.095623	29.15511449	-1.822162195	Down	0	0	AAEL001668-PA OS=Aedes aegypti GN=AAEL001668 PE=3 SV=1
CCG017822.1	2000	9	8598	0.19336381	150.2856318	9.602173597	Up	1.09E-06	2.01E-06	AAEL005922-PA OS=Aedes aegypti GN=AAEL005922 PE=4 SV=1
CCG009866.1	477	905	8608	81.5253875	630.8613133	2.952001584	Up	0	0	AAEL009985-PA OS=Aedes aegypti GN=AAEL009985 PE=4 SV=1
CCG006098.1	2403	2151	8616	38.4635452	125.3435342	1.704323929	Up	0	0	AAEL004704-PA OS=Aedes aegypti GN=AAEL004704 PE=4 SV=1
CCG008332.1	5517	2588	8643	20.1569101	54.766066	1.442007748	Up	0	0	AAEL008905-PA OS=Aedes aegypti GN=AAEL008905 PE=4 SV=1
CCG003404.1	468	26707	8683	2452.12116	648.5955496	-1.918639234	Down	0	0	AAEL010205-PA OS=Aedes aegypti GN=AAEL010205 PE=4 SV=1
CCG025782.1	14209	2845	8698	8.60362411	21.39961231	1.31456826	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018110 PE=4 SV=1
CCG027115.1	15692	2760	8704	7.55776622	19.39057106	1.359323493	Up	0	0	AAEL005356-PA OS=Aedes aegypti GN=AAEL005356 PE=4 SV=1
CCG027309.1	43656	3250	8710	3.19891052	6.974679098	1.124546205	Up	0	0	AAEL004125-PA OS=Aedes aegypti GN=AAEL004125 PE=4 SV=1
CCG005824.1	4173	4	8718	0.04118834	73.03289166	10.79209469	Up	2.95E-05	5.04E-05	AAEL000961-PA OS=Aedes aegypti GN=AAEL000961 PE=4 SV=1
CCG015500.2	30287	3007	8721	4.26618655	10.0660723	1.238481908	Up	0	0	AAEL014695-PA OS=Aedes aegypti GN=AAEL014695 PE=4 SV=1
CCG026155.1	213	2545	8725	513.417721	1431.976361	1.479802679	Up	0	0	Translation initiation factor 5A OS=Aedes albopictus PE=2 SV=1
CCG027401.1	4324	3554	8765	35.3178624	70.86246436	1.004623613	Up	1.85E-11	4.07E-11	DNA-directed RNA polymerase OS=Aedes aegypti GN=AAEL001847 PE=3 SV=1
CCG007724.1	1275	40703	8806	1371.76244	241.4451649	-2.506263179	Down	0	0	ATP synthase subunit beta OS=Anopheles darlingi GN=AND_21925 PE=3 SV=1
CCG023209.1	453	1313	8843	124.545831	682.4195177	2.453982177	Up	8.52E-13	2.13E-12	Putative salivary secreted peptide OS=Aedes albopictus PE=2 SV=1
CCG012536.1	1704	630	8883	15.8866977	182.2384816	3.519936461	Up	3.95E-13	1.05E-12	AAEL012817-PA OS=Aedes aegypti GN=AAEL012817 PE=3 SV=1

CCG025277.1	9296	22670	8905	104.789574	33.48789314	-1.645783671	Down	0	0	AAEL001668-PA OS=Aedes aegypti GN=AAEL001668 PE=3 SV=1
CCG012161.1	11696	3077	8919	11.3045377	26.65807717	1.237670724	Up	0	0	AAEL014508-PA OS=Aedes aegypti GN=AAEL014508 PE=4 SV=1
CCG025245.1	3717	486	8944	5.61831881	84.11805958	3.904205173	Up	0	0	AAEL014533-PA OS=Aedes aegypti GN=AAEL014533 PE=4 SV=1
CCG012408.1	7233	22297	8955	132.461796	43.28098555	-1.613771071	Down	0	0	AAEL006307-PA OS=Aedes aegypti GN=AAEL006307 PE=4 SV=1
CCG018796.1	3702	717	8995	8.3223392	84.94049151	3.351391461	Up	1.89E-14	6.62E-14	AAEL004017-PA OS=Aedes aegypti GN=AAEL004017 PE=4 SV=1
CCG023350.1	3282	43648	9002	571.46343	95.88495049	-2.575284876	Down	0	0	AAEL012897-PA OS=Aedes aegypti GN=AAEL012897 PE=4 SV=1
CCG018447.1	645	1324	9043	88.2045405	490.1204602	2.474211545	Up	1.05E-12	2.59E-12	AAEL009287-PA OS=Aedes aegypti GN=AAEL009287 PE=4 SV=1
CCG022464.1	13645	39632	9045	124.805901	23.17314866	-2.429160162	Down	0	0	AAEL008880-PA OS=Aedes aegypti GN=AAEL008880 PE=4 SV=1
CCG002334.1	1935	2898	9082	64.3546723	164.0780722	1.350265646	Up	0	0	Putative secreted peptide OS=Aedes albopictus PE=4 SV=1
CCG013504.1	2265	2418	9127	45.872326	140.867193	1.618639691	Up	0	0	AAEL001352-PA OS=Aedes aegypti GN=AAEL001352 PE=4 SV=1
CCG025494.1	4771	87831	9131	791.044804	66.90505664	-3.563572252	Down	0	0	Glyceroldehyde-3-phosphate dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005597 PE=3 SV=1
CCG022794.1	4000	28323	9163	304.257948	80.08067252	-1.925768957	Down	0	0	Sugar transporter ERD6-like 6 (Fragment) OS=Camponotus floridanus GN=EAG_03869 PE=3 SV=1
CCG016917.1	1559	1671	9174	46.05672	205.713426	2.159152378	Up	0	0	-
CCG016858.1	5025	700	9195	5.98583368	63.96842884	3.417735951	Up	0	0	AAEL003671-PA OS=Aedes aegypti GN=AAEL003671 PE=4 SV=1
CCG003573.1	396	2207	9221	239.480314	814.0158336	1.765149791	Up	0	0	AAEL002534-PA OS=Aedes aegypti GN=AAEL002534 PE=2 SV=1
CCG012680.1	6068	718	9249	5.08442147	53.28429497	3.389554845	Up	0	0	AAEL007252-PA OS=Aedes aegypti GN=AAEL007252 PE=4 SV=1

CCG003438.1	5332	1145	9261	9.22737174	60.71804223	2.718133589	Up	4.74E-12	1.08E-11	DNA topoisomerase 2 (Fragment) OS=Aedes aegypti GN=AAEL012584 PE=3 SV=1
CCG024769.1	2595	29015	9280	480.449653	125.0145697	-1.942289015	Down	0	0	AAEL010975-PA OS=Aedes aegypti GN=AAEL010975 PE=4 SV=1
CCG023696.1	5153	3771	9284	31.4455403	62.98324113	1.002112576	Up	4.81E-12	1.10E-11	AAEL004227-PA (Fragment) OS=Aedes aegypti GN=AAEL004227 PE=4 SV=1
CCG016749.1	2358	3124	9325	56.9285203	138.24679	1.280022476	Up	0	0	Forkhead box transcription factor subgroup K2 OS=Aedes aegypti PE=2 SV=1
CCG005596.1	4166	2681	9341	27.6528705	78.38340454	1.50311899	Up	0	0	AAEL008686-PA OS=Aedes aegypti GN=AAEL008686 PE=4 SV=1
CCG028566.1	5307	1718	9349	13.9103079	61.58374403	2.146395225	Up	2.10E-12	4.99E-12	AAEL004320-PA OS=Aedes aegypti GN=AAEL004320 PE=4 SV=1
CCG018893.1	1071	889	9362	35.6676882	305.5829945	3.098874645	Up	0	0	Histone H3 OS=Culex quinquefasciatus GN=CpipJ_CPIJ017395 PE=3 SV=1
CCG014460.1	3588	993	9390	11.8921255	91.48779791	2.943572739	Up	4.50E-12	1.04E-11	AAEL013285-PA OS=Aedes aegypti GN=AAEL013285 PE=4 SV=1
CCG007810.2	1560	65608	9458	1807.15279	211.9457575	-3.091951492	Down	0	0	AAEL007306-PA OS=Aedes aegypti GN=AAEL007306 PE=4 SV=1
CCG017994.1	24512	15627	9462	27.3942576	13.49441966	-1.021510573	Down	0	0	AAEL003046-PA OS=Aedes aegypti GN=AAEL003046 PE=4 SV=1
CCG024902.1	16796	1579	9523	4.03960532	19.82065192	2.294718165	Up	4.25E-12	9.79E-12	AAEL000172-PA OS=Aedes aegypti GN=AAEL000172 PE=4 SV=1
CCG021676.2	11172	29112	9527	111.970544	29.81091146	-1.909206858	Down	0	0	AAEL003078-PA OS=Aedes aegypti GN=AAEL003078 PE=4 SV=1
CCG010428.1	24232	3191	9528	5.6584856	13.74556211	1.280478008	Up	0	0	AAEL001141-PA OS=Aedes aegypti GN=AAEL001141 PE=4 SV=1
CCG000136.1	3773	1152	9562	13.1198342	88.59555857	2.755484893	Up	4.48E-12	1.03E-11	AAEL002407-PA OS=Aedes aegypti GN=AAEL002407 PE=4 SV=1

CCG014779.1	4131	21998	9573	228.818258	81.01079243	-1.498016145	Down	0	0	Anaphne ubiquitin-conjugating enzyme E2 binding protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005416 PE=4 SV=1
CCG014650.1	435	1338	9576	132.168977	769.5642721	2.54165824	Up	3.27E-12	7.64E-12	CEC-A OS=Anopheles melas GN=CEC-A PE=4 SV=1
CCG026177.1	1362	1841	9639	58.0817044	247.4029587	2.090707059	Up	0	0	AAEL001623-PA OS=Aedes aegypti GN=AAEL001623 PE=2 SV=1
CCG020090.1	6793	1646	9685	10.4119216	49.84114685	2.259100917	Up	3.63E-12	8.44E-12	AAEL001372-PA OS=Aedes aegypti GN=AAEL001372 PE=4 SV=1
CCG027469.1	2841	1895	9693	28.6616146	119.2715863	2.057058607	Up	0	0	AAEL000565-PA OS=Aedes aegypti GN=AAEL000565 PE=4 SV=1
CCG013234.1	869	2251	9699	111.305952	390.1729878	1.809583155	Up	0	0	Ribosomal protein S19 OS=Aedes albopictus PE=2 SV=1
CCG008472.1	3916	3084	9699	33.8403119	86.58333157	1.355346447	Up	0	0	AAEL004358-PA OS=Aedes aegypti GN=AAEL004358 PE=4 SV=1
CCG022635.1	6728	2679	9719	17.1099761	50.49932996	1.561426503	Up	0	0	Ubiquitin carboxyl-terminal hydrolase (Fragment) OS=Aedes aegypti GN=AAEL009824 PE=3 SV=1
CCG022647.1	1641	881	9722	23.0690653	207.1080846	3.166352414	Up	4.13E-13	1.09E-12	AAEL006389-PA (Fragment) OS=Aedes aegypti GN=AAEL006389 PE=3 SV=1
CCG026483.2	2090	3886	9741	79.8949228	162.9323321	1.028097188	Up	7.49E-12	1.68E-11	AAEL009646-PA OS=Aedes aegypti GN=AAEL009646 PE=4 SV=1
CCG022035.1	1095	16390	9748	643.172557	311.2084767	-1.047324494	Down	0	0	Fructose-bisphosphate aldolase OS=Aedes aegypti GN=AAEL005766 PE=3 SV=1
CCG010459.3	4605	63184	9770	589.576485	74.16772292	-2.990815602	Down	0	0	AAEL010975-PA OS=Aedes aegypti GN=AAEL010975 PE=4 SV=1
CCG018749.1	1473	16161	9776	471.441874	232.0109394	-1.022887077	Down	0	0	AAEL006866-PA OS=Aedes aegypti GN=AAEL006866 PE=4 SV=1
CCG005499.1	2661	326	9792	5.26423656	128.6401526	4.610972892	Up	6.17E-13	1.58E-12	AAEL013692-PA OS=Aedes aegypti GN=PIWI3 PE=4 SV=1

CCG020968.4	5553	1047	9802	8.10180301	61.7073706	2.929127908	Up	5.84E-12	1.32E-11	AAEL009381-PA OS=Aedes aegypti GN=AAEL009381 PE=4 SV=1
CCG024888.1	1545	3039	9894	84.5210508	223.8687316	1.405270428	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018253 PE=4 SV=1
CCG027993.1	692	1304	9919	80.9718698	501.085473	2.629564015	Up	4.00E-12	9.24E-12	AAEL000425-PA OS=Aedes aegypti GN=AAEL006425 PE=3 SV=1
CCG017862.1	852	3249	9979	163.859939	409.4467652	1.32121271	Up	0	0	AAEL000953-PA OS=Aedes aegypti GN=AAEL006953 PE=4 SV=1
CCG018906.1	2758	391	9981	6.09179341	126.5114432	4.376257058	Up	9.02E-13	2.25E-12	AAEL009508-PA OS=Aedes aegypti GN=AAEL009508 PE=4 SV=1
CCG026342.2	3129	2900	10004	39.8249378	111.768169	1.488765361	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ007196 PE=3 SV=1
CCG012974.1	4996	1677	10012	14.4235879	70.05649861	2.280088806	Up	3.09E-12	7.24E-12	Sub1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ013695 PE=4 SV=1
CCG012635.1	3786	356	10073	4.04047161	93.0096994	4.524785572	Up	7.93E-13	1.99E-12	AAEL013510-PA OS=Aedes aegypti GN=AAEL013510 PE=4 SV=1
CCG024907.1	1628	3739	10176	98.687861	218.510703	1.146759403	Up	1.04E-11	2.31E-11	AAEL013675-PA OS=Aedes aegypti GN=AAEL013675 PE=4 SV=1
CCG022191.1	2333	946	10199	17.4236472	152.8244598	3.132756915	Up	0	0	AAEL007880-PA OS=Aedes aegypti GN=AAEL007880 PE=3 SV=1
CCG022898.1	4878	2206	10428	19.4323974	74.73245391	1.943270996	Up	0	0	AAEL009073-PA OS=Aedes aegypti GN=AAEL009073 PE=4 SV=1
CCG020472.1	2469	3972	10436	69.1274953	147.7620804	1.095944525	Up	1.15E-11	2.55E-11	AAEL001432-PA OS=Aedes aegypti GN=AAEL001432 PE=3 SV=1
CCG023451.1	1615	119939	10439	3191.17461	225.9625085	-3.819932224	Down	0	0	Glyceroldehyde-3-phosphate dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ005597 PE=3 SV=1
CCG022045.1	2019	225	10452	4.78860341	180.9727136	5.240023422	Up	0	0	AAEL009903-PA (Fragment) OS=Aedes aegypti GN=AAEL009903 PE=4 SV=1

CCG002661.1	1266	2040	10468	69.2403307	289.0546929	2.061657976	Up	0	0	ATP-dependent RNA helicase WM6 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014241 PE=4 SV=1
CCG013870.1	498	903	10519	77.9150007	738.4058501	3.244440966	Up	0	0	AAEL0006860-PA OS=Aedes aegypti GN=AAEL006860 PE=4 SV=1
CCG013150.1	6030	2018	10530	14.3802528	61.04654302	2.085820561	Up	0	0	AAEL013136-PA OS=Aedes aegypti GN=AAEL013136 PE=4 SV=1
CCG013172.1	1749	34354	10612	844.015016	212.108195	-1.992468304	Down	0	0	AAEL000667-PA OS=Aedes aegypti GN=AAEL000667 PE=4 SV=1
CCG000824.1	6669	3093	10635	19.9288333	55.74767933	1.484054499	Up	0	0	AAEL007181-PA (Fragment) OS=Aedes aegypti GN=AAEL007181 PE=4 SV=1
CCG006267.1	7587	2371	10637	13.428396	49.01162383	1.867836954	Up	0	0	AAEL008484-PA OS=Aedes aegypti GN=AAEL008484 PE=4 SV=1
CCG011098.1	2217	3806	10638	73.7676184	167.7429627	1.185192679	Up	4.12E-12	9.51E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002058 PE=4 SV=1
CCG001421.1	1791	19136	10716	459.111581	209.1640949	-1.134209605	Down	0	0	AAEL005849-PA OS=Aedes aegypti GN=AAEL005849 PE=4 SV=1
CCG000166.1	3090	1440	10746	20.0247307	121.573347	2.60197223	Up	1.98E-12	4.73E-12	AAEL002551-PA OS=Aedes aegypti GN=AAEL002551 PE=4 SV=1
CCG008656.1	6387	2181	10840	14.6730846	59.33109758	2.015616285	Up	0	0	AAEL006038-PA (Fragment) OS=Aedes aegypti GN=AAEL006038 PE=4 SV=1
CCG014379.1	4203	58	10847	0.59296802	90.21946899	7.249340674	Up	0	0	AAEL011312-PA OS=Aedes aegypti GN=AAEL011312 PE=4 SV=1
CCG004880.1	6171	3660	10872	25.4852096	61.58910794	1.273015011	Up	7.23E-13	1.82E-12	AAEL005861-PA OS=Aedes aegypti GN=AAEL005861 PE=4 SV=1
CCG009668.1	2039	884	10911	18.6293504	187.067071	3.327906356	Up	0	0	AAEL012183-PA OS=Aedes aegypti GN=AAEL012183 PE=4 SV=1
CCG001360.1	28158	1781	10946	2.71784563	13.58950556	2.321957544	Up	0	0	AAEL001440-PA OS=Aedes aegypti GN=AAEL001440 PE=4 SV=1
CCG021834.1	2493	2610	11008	44.9863648	154.3604937	1.778743867	Up	0	0	AGAP011566-PA (Fragment) OS=Anopheles gambiae GN=AGAP011566 PE=4 SV=4

CCG020548.1	2682	2874	11055	46.0458677	144.0953578	1.645880262	Up	0	0	AAEL012930-PB (Fragment) OS=Aedes aegypti GN=AAEL012930 PE=4 SV=1
CCG013265.2	2271	2610	11111	49.383975	171.0354087	1.792180158	Up	0	0	AAEL004699-PA OS=Aedes aegypti GN=AAEL004699 PE=4 SV=1
CCG021922.2	8001	2853	11185	15.3221663	48.86993197	1.673326808	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006929 PE=4 SV=1
CCG024865.1	1759	56681	11199	1384.63191	222.5683579	-2.637182093	Down	0	0	AAEL005435-PA OS=Aedes aegypti GN=AAEL005435 PE=3 SV=1
CCG024101.1	5511	111	11265	0.86547642	71.4579909	6.367457031	Up	0	0	AAEL004462-PA OS=Aedes aegypti GN=AAEL004462 PE=4 SV=1
CCG025470.1	7088	3273	11367	19.8419782	56.0624622	1.498479242	Up	0	0	Rho guanine nucleotide exchange factor OS=Culex quinquefasciatus GN=CpipJ_CPIJ015585 PE=4 SV=1
CCG013880.1	24238	4241	11401	7.51855122	16.44357263	1.128997187	Up	2.01E-11	4.41E-11	AAEL001421-PA OS=Aedes aegypti GN=AAEL001421 PE=4 SV=1
CCG020798.1	1776	1044	11444	25.2592359	225.2604276	3.15671099	Up	2.08E-12	4.96E-12	AAEL009487-PA OS=Aedes aegypti GN=AAEL009487 PE=4 SV=1
CCG004583.4	7200	1403	11479	8.37313023	55.73417487	2.734723256	Up	1.72E-12	4.12E-12	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ010152 PE=4 SV=1
CCG012162.1	16917	1372	11496	3.48492498	23.75600578	2.769092784	Up	6.08E-12	1.37E-11	Mitotin OS=Culex quinquefasciatus GN=CpipJ_CPIJ010145 PE=4 SV=1
CCG003361.1	4635	1353	11518	12.5432688	86.87150634	2.791969687	Up	2.77E-12	6.50E-12	AAEL005567-PA OS=Aedes aegypti GN=AAEL005567 PE=2 SV=1
CCG021110.1	822	3715	11523	194.2002	490.0537996	1.335395458	Up	2.75E-12	6.47E-12	AAEL012359-PA OS=Aedes aegypti GN=AAEL012359 PE=3 SV=1
CCG014982.1	4080	3804	11536	40.0629584	98.84281393	1.302867122	Up	3.42E-12	7.96E-12	AAEL006956-PA OS=Aedes aegypti GN=AAEL006956 PE=4 SV=1
CCG017228.1	16861	3443	11547	8.77437852	23.94064539	1.448093195	Up	1.89E-11	4.16E-11	AAEL007841-PA OS=Aedes aegypti GN=AAEL007841 PE=4 SV=1
CCG018955.1	6207	2394	11642	16.5731504	65.56859323	1.984157069	Up	0	0	AAEL005712-PA OS=Aedes aegypti GN=AAEL005712 PE=4 SV=1

CCG015850.1	13899	1357	11690	4.1952608	29.40227754	2.809095508	Up	3.94E-12	9.11E-12	AAEL000416-PA OS=Aedes aegypti GN=AAEL000416 PE=4 SV=1
CCG002028.1	9225	2084	11695	9.70720075	44.31837906	2.190777883	Up	0	0	AAEL009884-PA OS=Aedes aegypti GN=AAEL009884 PE=4 SV=1
CCG006928.1	48411	4644	11703	4.12202697	8.45090399	1.035751731	Up	1.24E-11	2.75E-11	Noncatalytic domain interacting protein kinase, isoform A OS=Drosophila melanogaster GN=hipk PE=4 SV=2
CCG008422.1	687	51611	11783	3228.10913	599.5827878	-2.428658478	Down	0	0	Mitochondrial ATP synthase lipid binding protein OS=Aedes albopictus PE=2 SV=1
CCG019250.1	2043	672	11799	14.1339509	201.8955985	3.836372753	Up	0	0	AAEL012817-PA OS=Aedes aegypti GN=AAEL012817 PE=3 SV=1
CCG003755.1	3590	1367	11818	16.3620132	115.0799206	2.81421396	Up	1.86E-12	4.45E-12	AAEL001037-PA OS=Aedes aegypti GN=AAEL001037 PE=3 SV=1
CCG009771.2	28964	60615	11859	89.9257859	14.31329251	-2.651379287	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_00717 PE=4 SV=1
CCG018238.1	2033	59232	11883	1251.93474	204.3331052	-2.615164489	Down	0	0	mitochondrial cytochrome c1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ014410 PE=4 SV=1
CCG018262.1	432	75481	11947	7507.86699	966.774381	-2.957151943	Down	0	0	AAEL009201-PA OS=Aedes aegypti GN=AAEL009201 PE=4 SV=1
CCG012826.1	1180	167776	11965	6109.56796	354.4710013	-4.10733085	Down	0	0	AAEL009185-PA OS=Aedes aegypti GN=AAEL009185 PE=2 SV=1
CCG011460.1	13700	158705	11981	497.774579	30.57190613	-4.025214144	Down	0	0	AAEL010850-PA OS=Aedes aegypti GN=AAEL010850 PE=4 SV=1
CCG025582.1	1830	2329	11987	54.6866185	228.9862643	2.066001298	Up	0	0	AAEL011255-PA OS=Aedes aegypti GN=AAEL011255 PE=4 SV=1
CCG021851.3	9276	1703	12097	7.88890233	45.58972338	2.530812174	Up	5.00E-12	1.14E-11	Hyperplastic discs protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011645 PE=4 SV=1
CCG014112.1	7421	3184	12131	18.4362802	57.14578298	1.632099444	Up	0	0	AGAP011241-PA (Fragment) OS=Anopheles gambiae GN=AGAP011241 PE=4 SV=3

CCG006846.1	4269	2229	12136	22.4360596	99.3801	2.147137673	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_17516 PE=4 SV=1
CCG026871.2	789	3333	12194	181.518537	540.2803891	1.573591432	Up	9.99E-12	2.22E-11	40S ribosomal protein S4 OS=Aedes aegypti PE=2 SV=1
CCG018112.1	1243	2639	12208	91.2285838	343.3392139	1.912076815	Up	0	0	AAEL003750-PA OS=Aedes aegypti GN=AAEL003750 PE=4 SV=1
CCG011606.1	42801	3741	12218	3.75574816	9.979211364	1.409825478	Up	6.83E-12	1.53E-11	AAEL009400-PB OS=Aedes aegypti GN=AAEL009400 PE=4 SV=1
CCG026208.1	95960	3527	12273	1.57934821	4.471060138	1.501287666	Up	1.40E-11	3.09E-11	AAEL007151-PA OS=Aedes aegypti GN=AAEL007151 PE=4 SV=1
CCG015186.1	24333	51538	12281	91.0111446	17.64363609	-2.36689531	Down	0	0	AGAP001633-PA OS=Anopheles gambiae GN=AgaP_AGAP001633 PE=4 SV=5
CCG016461.1	3461	1439	12323	17.8657752	124.4700505	2.800528226	Up	3.28E-12	7.67E-12	GE12944 OS=Drosophila yakuba GN=GE12944 PE=4 SV=1
CCG014657.1	1968	2449	12331	53.4719919	219.0398937	2.034338327	Up	0	0	PolyQ domain-containing hypothetical conserved protein OS=Aedes aegypti PE=2 SV=1
CCG013859.1	58746	3297	12339	2.41158913	7.342630598	1.606312889	Up	0	0	AAEL006783-PA OS=Aedes aegypti GN=AAEL006783 PE=4 SV=1
CCG008569.1	1483	2186	12367	63.3390694	291.5232696	2.20244347	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_10459 PE=3 SV=1
CCG021882.1	4137	4928	12372	51.185606	104.5452744	1.030317781	Up	1.78E-11	3.91E-11	AAEL012243-PA OS=Aedes aegypti GN=AAEL012243 PE=4 SV=1
CCG018594.1	4923	2644	12391	23.0777937	87.98862634	1.930811745	Up	0	0	AAEL009189-PA OS=Aedes aegypti GN=AAEL009189 PE=4 SV=1
CCG005068.1	4719	27398	12400	249.477599	91.85900233	-1.441417259	Down	0	0	AAEL002879-PA (Fragment) OS=Aedes aegypti GN=AAEL002879 PE=4 SV=1
CCG011256.1	447	4777	12482	459.208998	976.1727308	1.087985539	Up	2.89E-11	6.28E-11	Ribosomal protein S16 OS=Aedes albopictus PE=2 SV=1
CCG000359.1	2235	466	12533	8.95923773	196.0322518	4.45157123	Up	0	0	AAEL007799-PA OS=Aedes aegypti GN=AAEL007799 PE=4 SV=1

CCG014445.1	20131	2726	12596	5.81866259	21.8734516	1.910421399	Up	0	0	AAEL014969-PA (Fragment) OS=Aedes aegypti GN=AAEL014969 PE=4 SV=1
CCG020936.1	936	3912	12606	179.591455	470.8162788	1.390445496	Up	2.06E-11	4.51E-11	AAEL013069-PA OS=Aedes aegypti GN=AAEL013069 PE=2 SV=1
CCG017214.1	7413	1866	12773	10.8163395	60.23500192	2.477389724	Up	0	0	AAEL001440-PA OS=Aedes aegypti GN=AAEL001440 PE=4 SV=1
CCG027737.1	10104	4167	12823	17.7211881	44.36559611	1.323966025	Up	3.13E-11	6.80E-11	Hypertrophic discs protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ011645 PE=4 SV=1
CCG027733.1	6999	1204	12839	7.39185033	64.12763473	3.116938745	Up	2.65E-13	7.40E-13	AAEL009884-PA OS=Aedes aegypti GN=AAEL009884 PE=4 SV=1
CCG021036.1	2285	3388	12856	63.7117991	196.6842915	1.626249258	Up	1.69E-11	3.72E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_14433 PE=4 SV=1
CCG004084.1	3492	84519	12880	1040.02262	128.9411812	-3.011829901	Down	0	0	myosin heavy chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ000853 PE=4 SV=1
CCG020899.1	876	4412	12961	216.418344	517.2308507	1.256985541	Up	1.12E-11	2.49E-11	AAEL013533-PA OS=Aedes aegypti GN=AAEL013533 PE=4 SV=1
CCG022449.1	1734	7	13019	0.1734649	262.4693225	10.5632893	Up	4.31E-06	7.70E-06	AAEL000776-PA OS=Aedes aegypti GN=AAEL000776 PE=4 SV=1
CCG008523.1	6612	0	13020	0.001	68.83798602	16.07091727	Up	4.44E-16	1.65E-15	AAEL008829-PA OS=Aedes aegypti GN=AAEL008829 PE=4 SV=1
CCG002280.1	604	22185	13053	1578.28404	755.4807727	-1.062889927	Down	0	0	Putative uncharacterized protein (Fragment) OS=Culex quinquefasciatus GN=CpipJ_CPIJ015860 PE=4 SV=1
CCG002188.1	558	4540	13081	349.610386	819.5147284	1.229021836	Up	2.32E-11	5.06E-11	60S ribosomal protein L27a OS=Aedes albopictus PE=2 SV=1
CCG002680.1	982	2485	13087	108.737058	465.8848963	2.099129866	Up	0	0	Ribosomal protein S19 OS=Aedes albopictus PE=2 SV=1
CCG019236.1	3366	1499	13127	19.1359573	136.3331251	2.832778161	Up	1.93E-12	4.62E-12	AAEL005813-PA OS=Aedes aegypti GN=AAEL005813 PE=4 SV=1

CCG003796.1	5786	879	13221	6.52789435	79.87960201	3.613137531	Up	0	0	AAEL001939-PA OS=Aedes aegypti GN=AAEL001939 PE=4 SV=1
CCG005343.1	3975	1834	13302	19.8255329	116.9849051	2.560890836	Up	0	0	Cupis ta nonucleoprotein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008950 PE=4 SV=1
CCG009349.1	2520	4034	13333	68.7856783	184.9598033	1.427031646	Up	2.35E-11	5.12E-11	AGAP001546-PA OS=Anopheles gambiae GN=AgaP_AGAP001546 PE=3 SV=4
CCG019635.1	4698	3300	13336	30.1830831	99.2344783	1.717101284	Up	0	0	AAEL014034-PA OS=Aedes aegypti GN=AAEL014034 PE=4 SV=1
CCG025338.1	747	24996	13453	1437.8467	629.5765695	-1.191456114	Down	0	0	14-3-3 zeta OS=Culex pipiens PE=2 SV=1
CCG023524.1	4298	4711	13533	47.0987482	110.0722102	1.224689659	Up	1.20E-11	2.65E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_09747 PE=4 SV=1
CCG021617.1	3497	2520	13652	30.9647501	136.4742335	2.139929885	Up	0	0	AAEL002370-PA OS=Aedes aegypti GN=AAEL002370 PE=4 SV=1
CCG025005.1	2519	3199	13714	54.5693454	190.3206858	1.802269734	Up	0	0	AAEL011704-PA OS=Aedes aegypti GN=AAEL011704 PE=3 SV=1
CCG010286.1	3675	1536	13780	17.9595952	131.0816473	2.867638971	Up	2.61E-12	6.14E-12	AAEL014711-PA OS=Aedes aegypti GN=Aael_AAEL014711 PE=4 SV=1
CCG002118.1	5345	35805	13823	287.844967	90.40753224	-1.670777106	Down	0	0	Ubiquitin C variant (Fragment) OS=Homo sapiens PE=2 SV=1
CCG019619.1	1769	119868	13841	2911.64282	273.5203555	-3.412113279	Down	0	0	ATP synthase subunit beta OS=Aedes aegypti GN=AAEL003393 PE=3 SV=1
CCG000263.3	35854	3623	13873	4.34203571	13.52641752	1.639336293	Up	7.40E-12	1.66E-11	AAEL004381-PA OS=Aedes aegypti GN=AAEL004381 PE=4 SV=1
CCG011939.1	3028	2924	13917	41.4938917	160.6718421	1.953146239	Up	0	0	AAEL012827-PA OS=Aedes aegypti GN=AAEL012827 PE=3 SV=1
CCG018919.1	969	2	13919	0.08868882	502.1509333	12.46708115	Up	9.36E-05	0.00015347	AAEL000923-PA OS=Aedes aegypti GN=AAEL000923 PE=4 SV=1
CCG018800.1	3708	1657	13951	19.2019553	131.5272166	2.776036233	Up	2.91E-12	6.82E-12	AAEL004035-PA OS=Aedes aegypti GN=AAEL004035 PE=4 SV=1

CCG010188.1	2130	3288	13993	66.3307453	229.6578248	1.791736299	Up	5.95E-12	1.34E-11	AAEL001352-PA OS=Aedes aegypti GN=AAEL001352 PE=4 SV=1
CCG027392.1	1728	23724	13996	589.938648	283.1458575	-1.059019503	Down	0	0	AAEL002833-PA OS=Aedes aegypti GN=AAEL002833 PE=3 SV=1
CCG022750.1	1920	13	13998	0.29094091	254.8676866	9.774806578	Up	6.34E-08	1.23E-07	AAEL002652-PA OS=Aedes aegypti GN=AAEL002652 PE=4 SV=1
CCG008471.1	5263	1165	14079	9.51163611	93.51654545	3.29745621	Up	1.90E-12	4.56E-12	AAEL004356-PA OS=Aedes aegypti GN=AAEL004356 PE=4 SV=1
CCG027950.1	13592	1280	14202	4.04659066	36.52718119	3.1741916	Up	2.19E-12	5.20E-12	AGAP005719-PA OS=Anopheles gambiae GN=AGAP005719 PE=4 SV=4
CCG014060.1	4620	2196	14262	20.4245752	107.9166544	2.401539553	Up	0	0	AAEL008027-PA OS=Aedes aegypti GN=AAEL008027 PE=4 SV=1
CCG001272.1	5666	2516	14262	19.080807	87.99416578	2.205285686	Up	0	0	AAEL000604-PA OS=Aedes aegypti GN=AAEL000604 PE=4 SV=1
CCG016448.1	1579	909	14295	24.7368517	316.4842093	3.677399719	Up	2.27E-12	5.39E-12	Basic helix-loop-helix transcription factor OS=Culex quinquefasciatus GN=CpipJ_CPIJ008846 PE=4 SV=1
CCG018567.2	4347	3916	14304	38.7093353	115.03179	1.571279175	Up	7.60E-12	1.70E-11	Helix-loop-helix transcription factor OS=Culex quinquefasciatus GN=CpipJ_CPIJ009239 PE=4 SV=1
CCG002354.1	585	4227	14331	310.483877	856.3881443	1.463746469	Up	1.15E-11	2.55E-11	40S ribosomal protein S17 OS=Aedes aegypti GN=AAEL004175 PE=2 SV=2
CCG000933.1	6377	961	14420	6.47544534	79.04945132	3.609704127	Up	1.90E-12	4.55E-12	AAEL002455-PA OS=Aedes aegypti GN=AAEL002455 PE=4 SV=1
CCG001877.1	4150	1716	14492	17.7677264	122.0759872	2.780448457	Up	1.20E-14	4.28E-14	Ubiquitin carboxyl-terminal hydrolase OS=Aedes aegypti GN=AAEL002698 PE=3 SV=1
CCG003480.1	4128	2699	14575	28.094795	123.4294777	2.135312205	Up	0	0	AAEL011704-PA OS=Aedes aegypti GN=AAEL011704 PE=3 SV=1
CCG012249.1	3181	92252	14638	1246.16283	160.8674176	-2.953548534	Down	0	0	AAEL012897-PA OS=Aedes aegypti GN=AAEL012897 PE=4 SV=1

CCG022705.1	4995	1385	14737	11.914531	103.1391641	3.113798188	Up	7.34E-12	1.64E-11	AAEL012568-PA OS=Aedes aegypti GN=AAEL012568 PE=4 SV=1
CCG006686.1	3507	5631	14758	68.9941761	147.1098513	1.092347369	Up	1.24E-11	2.75E-11	AAEL003976-PA OS=Aedes aegypti GN=AAEL003976 PE=4 SV=1
CCG003600.1	7932	2430	14772	13.1639505	65.10384071	2.306150153	Up	0	0	AAEL009391-PA OS=Aedes aegypti GN=AAEL009391 PE=4 SV=1
CCG007312.1	24630	5257	14778	9.1714127	20.97496606	1.193452597	Up	4.30E-11	9.28E-11	Cinfefer OS=Culex quinquefasciatus GN=CpipJ_CPIJ009239 PE=4 SV=1
CCG009037.1	32742	1411	14818	1.85175907	15.8210172	3.09487405	Up	6.09E-12	1.37E-11	AGAP012403-PA (Fragment) OS=Anopheles gambiae GN=AGAP012403 PE=4 SV=4
CCG014351.1	11145	3050	14824	11.7593262	46.49811529	1.983366843	Up	0	0	AAEL012446-PA OS=Aedes aegypti GN=IAP6 PE=4 SV=2
CCG016498.1	3826	5913	14839	66.4087927	135.5843881	1.029744894	Up	0	0	AGAP010331-PA OS=Anopheles gambiae GN=AGAP010331 PE=3 SV=3
CCG008592.1	3446	4565	14859	56.9230524	150.7385475	1.404963465	Up	1.55E-11	3.42E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_02441 PE=3 SV=1
CCG025588.1	60860	3336	14907	2.35535713	8.562652511	1.862111958	Up	0	0	AAEL002435-PA OS=Aedes aegypti GN=AAEL002435 PE=4 SV=1
CCG019910.1	3690	2389	15021	27.8197008	142.3057657	2.354815306	Up	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_00689 PE=4 SV=1
CCG018886.1	339	5581	15090	707.416191	1556.107364	1.137310458	Up	6.47E-12	1.46E-11	AAEL011414-PA OS=Aedes aegypti GN=AAEL011414 PE=4 SV=1
CCG012846.1	9297	3735	15116	17.2627685	56.83869116	1.719209479	Up	1.99E-11	4.35E-11	AAEL000172-PA OS=Aedes aegypti GN=AAEL000172 PE=4 SV=1
CCG001809.1	6930	1368	15141	8.48233722	76.37853804	3.170633561	Up	4.04E-12	9.34E-12	AAEL003882-PA OS=Aedes aegypti GN=AAEL003882 PE=4 SV=1
CCG013403.1	5103	4565	15144	38.4395137	103.7444921	1.432372755	Up	1.55E-11	3.42E-11	Protein kinase OS=Culex quinquefasciatus GN=CpipJ_CPIJ014503 PE=4 SV=1

CCG017490.1	17590	5760	15193	14.070817	30.19449113	1.101579259	Up	2.89E-11	6.28E-11	Nuclear receptor co-repressor 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002507 PE=4 SV=1
CCG005737.1	1189	1080	15310	39.0305411	450.1355907	3.527684269	Up	6.28E-12	1.41E-11	AAEL000823-PA OS=Aedes aegypti GN=AAEL000823 PE=4 SV=1
CCG013157.1	31843	4345	15325	5.86325085	16.82428137	1.520772196	Up	2.71E-12	6.38E-12	AGAP003043-PA OS=Anopheles gambiae GN=AgaP_AGAP003043 PE=4 SV=5
CCG005186.1	3885	3590	15431	39.7069105	138.8522957	1.806089013	Up	7.25E-12	1.62E-11	AAEL003126-PA OS=Aedes aegypti GN=AAEL003126 PE=4 SV=1
CCG016139.1	586	4661	15527	341.778043	926.2750229	1.438380897	Up	1.41E-11	3.11E-11	40S ribosomal protein S17 OS=Aedes aegypti GN=AAEL004175 PE=2 SV=2
CCG006475.1	369	4317	15557	502.71096	1473.837159	1.551776085	Up	1.51E-11	3.33E-11	40S ribosomal protein S21 OS=Aedes aegypti GN=AAEL000529 PE=3 SV=1
CCG027712.1	1076	6103	15716	243.721459	510.5987711	1.066956781	Up	6.56E-12	1.47E-11	AAEL011711-PA OS=Aedes aegypti GN=AAEL011711 PE=4 SV=1
CCG026788.1	2058	4261	15749	88.9669772	267.5208459	1.588309473	Up	1.77E-11	3.90E-11	AAEL008958-PA OS=Aedes aegypti GN=FoxK1 PE=2 SV=1
CCG014615.1	543	3561	15770	281.795994	1015.270764	1.849141523	Up	1.92E-11	4.23E-11	CEC-A OS=Anopheles melas GN=CEC-A PE=4 SV=1
CCG016790.1	492	1805	15858	157.64303	1126.764945	2.837453292	Up	0	0	AAEL005722-PA OS=Aedes aegypti GN=AAEL005722 PE=4 SV=1
CCG004790.1	2449	3192	15884	56.0062854	226.7363284	2.017354913	Up	0	0	AAEL009955-PA OS=Aedes aegypti GN=AAEL009955 PE=4 SV=1
CCG026417.1	3048	3552	15896	50.0749663	182.3152125	1.864273495	Up	7.46E-12	1.67E-11	AAEL010312-PA OS=Aedes aegypti GN=AAEL010312 PE=4 SV=1
CCG015624.1	21671	4366	15987	8.65700066	25.78921004	1.574828407	Up	1.40E-11	3.10E-11	AAEL007841-PA OS=Aedes aegypti GN=AAEL007841 PE=4 SV=1
CCG022109.1	3789	574	16030	6.50953488	147.8968532	4.505893082	Up	3.10E-13	8.52E-13	AAEL013510-PA OS=Aedes aegypti GN=AAEL013510 PE=4 SV=1
CCG014981.1	1155	87782	16120	3265.77424	487.9025295	-2.742760193	Down	0	0	AAEL008844-PA OS=Aedes aegypti GN=AAEL008844 PE=4 SV=1

CCG008776.1	1392	745	16225	22.9974514	407.4698568	4.147147446	Up	0	0	AAEL014238-PA OS=Aedes aegypti GN=AAEL014238 PE=3 SV=1
CCG026834.1	3979	3108	16359	33.5636932	143.7251696	2.098339356	Up	0	0	AAEL001791-PA (Fragment) OS=Aedes aegypti GN=AAEL001791 PE=4 SV=1
CCG002544.1	913	3111	16365	146.417135	626.6070092	2.097476506	Up	0	0	60S ribosomal protein L11 OS=Aedes albopictus PE=2 SV=1
CCG008753.1	1158	3560	16484	132.100393	497.6271453	1.913430426	Up	1.18E-11	2.61E-11	60S ribosomal protein L2/L8 OS=Aedes aegypti GN=AAEL000987 PE=2 SV=1
CCG022670.1	1593	4428	16563	119.441296	363.4739097	1.605550084	Up	8.68E-12	1.94E-11	Putative 56 kDa salivary secreted protein OS=Aedes albopictus PE=2 SV=1
CCG009249.1	813	3200	16600	169.130567	713.785233	2.077352635	Up	0	0	40S ribosomal protein S3a (Fragment) OS=Aedes triseriatus PE=2 SV=1
CCG013173.1	5250	34632	16661	283.452924	110.9409237	-1.353317507	Down	0	0	Uncharacterized protein OS=Anopheles darlingi GN=AND_20280 PE=4 SV=1
CCG023200.1	13842	56992	16940	176.920323	42.78234421	-2.048012343	Down	0	0	AAEL005324-PA OS=Aedes aegypti GN=AAEL005324 PE=4 SV=1
CCG027241.1	3390	1310	17096	16.6048237	176.2969616	3.4083333	Up	2.61E-12	6.14E-12	Importin-7 OS=Culex quinquefasciatus GN=CpipJ_CPIJ002096 PE=4 SV=1
CCG026048.1	711	32500	17324	1964.15805	851.7822599	-1.205354437	Down	0	0	60S ribosomal protein L13a (Fragment) OS=Aedes aegypti PE=2 SV=1
CCG018788.1	930	6200	17370	286.464897	652.9303952	1.188570837	Up	0	0	AAEL009653-PA OS=Aedes aegypti GN=AAEL009653 PE=4 SV=1
CCG027185.1	9328	30266	17392	139.421311	65.17949717	-1.09696097	Down	0	0	AAEL007019-PA OS=Aedes aegypti GN=AAEL007019 PE=4 SV=1
CCG018795.1	807	1663	17476	88.5485361	757.0394608	3.095828141	Up	3.33E-12	7.77E-12	AAEL006820-PA OS=Aedes aegypti GN=AAEL006820 PE=4 SV=1

CCG014475.1	5517	48764	17835	379.803542	113.0108512	-1.748792052	Down	0	0	AGAP003649-PA OS=Anopheles gambiae GN=AgaP_AGAP003649 PE=4 SV=1
CCG005442.1	606	4101	17914	290.790234	1033.403581	1.829353066	Up	3.05E-11	6.63E-11	Ribosomal protein S5 OS=Aedes albopictus PE=2 SV=1
CCG013505.1	8618	5565	18103	27.7473397	73.43347464	1.404088409	Up	1.26E-11	2.79E-11	Thyroid hormone receptor interactor 12 OS=Culex quinquefasciatus GN=CqipJ_CPIJ013812 PE=4 SV=1
CCG019329.1	14154	5311	18249	16.1235171	45.07231827	1.483075179	Up	1.58E-11	3.49E-11	AAEL007132-PA OS=Aedes aegypti GN=AAEL007132 PE=4 SV=1
CCG015704.1	10974	3169	18442	12.4085191	58.74799893	2.243208761	Up	0	0	AAEL013136-PA OS=Aedes aegypti GN=AAEL013136 PE=4 SV=1
CCG008638.3	2804	2145	18444	32.8709275	229.9466679	2.806415222	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CqipJ_CPIJ003009 PE=4 SV=1
CCG014921.1	981	4	18505	0.17520789	659.432122	11.87794072	Up	2.95E-05	5.02E-05	AAEL000442-PA OS=Aedes aegypti GN=AAEL000442 PE=4 SV=1
CCG013291.2	2727	6110	18543	96.2761564	237.7085905	1.30394359	Up	3.97E-12	9.17E-12	AAEL011387-PA (Fragment) OS=Aedes aegypti GN=AAEL011387 PE=4 SV=1
CCG022771.1	27312	1533	18573	2.41185571	23.77270337	3.301090467	Up	5.99E-12	1.35E-11	AAEL003882-PA OS=Aedes aegypti GN=AAEL003882 PE=4 SV=1
CCG010187.2	9057	4738	18583	22.4788123	71.72680327	1.673946526	Up	2.77E-11	6.03E-11	Thyroid hormone receptor interactor 12 OS=Culex quinquefasciatus GN=CqipJ_CPIJ013812 PE=4 SV=1
CCG025714.1	2421	6524	18596	115.792874	268.5188413	1.213476848	Up	2.85E-11	6.19E-11	AAEL011704-PA OS=Aedes aegypti GN=AAEL011704 PE=3 SV=1
CCG013849.1	1973	553	18828	12.0437219	333.6008287	4.791769671	Up	7.29E-13	1.83E-12	AAEL010094-PA OS=Aedes aegypti GN=AAEL010094 PE=3 SV=1
CCG007885.1	3489	136725	18901	1683.87417	189.3798765	-3.152429388	Down	0	0	Phosphorylase OS=Aedes aegypti GN=AAEL000703 PE=3 SV=1
CCG019897.1	29638	3482	18997	5.04826965	22.40712544	2.150096702	Up	1.80E-11	3.95E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CqipJ_CPIJ002396 PE=4 SV=1

CCG023764.1	1164	979	19397	36.1403524	582.5478476	4.010694072	Up	2.06E-12	4.91E-12	Peptidyl-prolyl cis-trans isomerase OS=Culex quinquefasciatus GN=CpipJ_CPIJ009378 PE=4 SV=1
CCG025358.4	10698	7022	19662	28.2046622	64.25029332	1.187769381	Up	0	0	Chromodomain helicase DNA binding protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003506 PE=4 SV=1
CCG000253.1	6548	4298	20020	28.2046303	106.8822085	1.92201779	Up	0	0	AAEL007417-PA OS=Aedes aegypti GN=AAEL007417 PE=4 SV=1
CCG010133.1	3276	48702	20213	638.800981	215.6934214	-1.566384354	Down	0	0	Elongation factor 1-alpha OS=Culex quinquefasciatus GN=CpipJ_CPIJ009655 PE=3 SV=1
CCG005443.1	936	6240	20386	286.464897	761.3882802	1.410273997	Up	9.71E-12	2.16E-11	60S ribosomal protein L27 OS=Aedes aegypti GN=AAEL011587 PE=3 SV=1
CCG019944.1	411	4689	20808	490.231352	1769.85845	1.852099324	Up	1.17E-11	2.60E-11	60S ribosomal protein L27 OS=Aedes aegypti GN=AAEL011587 PE=3 SV=1
CCG021934.1	2134	103537	20814	2084.79729	340.9660603	-2.612207066	Down	0	0	ATP synthase subunit beta OS=Aedes aegypti GN=AAEL002827 PE=3 SV=1
CCG025223.1	832	5212	20968	269.180597	881.0158021	1.710593474	Up	1.95E-11	4.28E-11	Ribosomal protein L26 OS=Aedes albopictus PE=2 SV=1
CCG017609.3	18332	7481	21130	17.5352708	40.29393346	1.200302935	Up	0	0	AAEL005380-PA OS=Aedes aegypti GN=AAEL005380 PE=4 SV=1
CCG009781.1	1308	120724	21651	3965.96196	578.655696	-2.776893744	Down	0	0	AAEL006572-PA OS=Aedes aegypti GN=AAEL006572 PE=4 SV=1
CCG018689.1	7050	4214	21793	25.6843208	108.0632235	2.072915794	Up	1.83E-11	4.03E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002709 PE=4 SV=1
CCG007854.1	1266	3359	21867	114.008956	603.8172497	2.404964809	Up	1.30E-11	2.87E-11	Ribosomal protein S2 OS=Aedes albopictus PE=2 SV=1
CCG005414.1	8407	3249	21958	16.606241	91.3065114	2.45899221	Up	0	0	AAEL003780-PA OS=Aedes aegypti GN=AAEL003780 PE=4 SV=1

CCG004829.1	1812	48	22157	1.13827112	427.4671846	8.55282564	Up	5.35E-14	1.80E-13	AAEL010097-PA OS=Aedes aegypti GN=AAEL010097 PE=4 SV=1
CCG008758.2	5703	1299	22275	9.78742509	136.5414019	3.8022653	Up	7.92E-13	1.99E-12	AAEL006249-PA OS=Aedes aegypti GN=AAEL006249 PE=4 SV=1
CCG025396.1	1353	5970	22582	189.600381	583.4647495	1.621683633	Up	2.89E-11	6.29E-11	GL12416 OS=Drosophila persimilis GN=GL12416 PE=3 SV=1
CCG027283.1	4936	7280	22611	63.3751353	160.1380867	1.337327647	Up	0	0	Ubiquitin carboxyl-terminal hydrolase OS=Culex quinquefasciatus GN=CpipJ_CPIJ010988 PE=3 SV=1
CCG001497.1	1221	7861	22611	276.64626	647.3723146	1.226553247	Up	5.39E-13	1.39E-12	AAEL003203-PA OS=Aedes aegypti GN=AAEL003203 PE=3 SV=1
CCG005199.1	819	3256	22634	170.829616	966.1118882	2.499632167	Up	0	0	AAEL006425-PA OS=Aedes aegypti GN=AAEL006425 PE=3 SV=1
CCG007782.1	2946	7600	22667	110.851997	268.9746299	1.278835342	Up	9.81E-12	2.18E-11	AAEL008738-PA OS=Aedes aegypti GN=AAEL008738 PE=3 SV=1
CCG006238.1	3465	5960	23214	73.9104237	234.2053209	1.663924103	Up	2.26E-11	4.93E-11	Uncharacterized protein OS=Anopheles darlingi GN=AND_09747 PE=4 SV=1
CCG025735.1	5682	13	23503	0.09831161	144.6012635	10.52243078	Up	6.34E-08	1.23E-07	AAEL000961-PA OS=Aedes aegypti GN=AAEL000961 PE=4 SV=1
CCG016378.1	715	9295	23640	558.60655	1155.823305	1.049016469	Up	0	0	40S ribosomal protein S20 OS=Aedes albopictus PE=2 SV=1
CCG021917.1	641	6362	23720	426.479644	1293.619852	1.600864937	Up	0	0	AAEL002372-PB OS=Aedes aegypti GN=RpS11 PE=2 SV=1
CCG022267.1	5172	1959	23897	16.2756593	161.5231903	3.310953417	Up	0	0	AAEL014711-PA OS=Aedes aegypti GN=Aael_AAEL014711 PE=4 SV=1
CCG009594.1	1155	1902	24121	70.76055	730.0680468	3.367013772	Up	0	0	AAEL010576-PA OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG020485.1	1424	50708	24162	1530.13294	593.161435	-1.367160292	Down	0	0	-
CCG011345.1	789	9941	25081	541.396872	1111.265576	1.037445185	Up	4.74E-12	1.08E-11	40S ribosomal protein S4 OS=Aedes aegypti PE=2 SV=1
CCG025006.1	8032	2576	25165	13.78113	109.5275193	2.990527303	Up	0	0	AAEL011704-PA OS=Aedes aegypti GN=AAEL011704 PE=3 SV=1

CCG002238.1	835	1739	25604	89.4902617	1071.942178	3.58235258	Up	4.99E-12	1.13E-11	Ribosomal protein L26 OS=Aedes albopictus PE=2 SV=1
CCG014731.1	35249	3487	25605	4.25077207	25.3938176	2.578680494	Up	1.24E-11	2.74E-11	AAEL001147-PA OS=Aedes aegypti GN=AAEL001147 PE=4 SV=1
CCG027724.1	447	6502	25829	625.0318	2019.994029	1.692349532	Up	2.60E-11	5.67E-11	60S ribosomal protein L18a OS=Aedes albopictus PE=2 SV=1
CCG007059.1	1143	9081	26027	341.389466	796.0271825	1.221399152	Up	1.58E-11	3.49E-11	Ribosomal protein L23a OS=Aedes albopictus PE=2 SV=1
CCG006597.1	710	9839	26095	595.463689	1284.839764	1.109503008	Up	0	0	Ribosomal protein L24 OS=Aedes albopictus PE=2 SV=1
CCG002564.1	978	5760	26258	253.073284	938.5832633	1.890929521	Up	2.89E-11	6.28E-11	Ribosomal protein L7AE OS=Aedes aegypti PE=2 SV=1
CCG007855.1	4890	7181	26365	63.1013628	188.4815884	1.578680533	Up	8.16E-12	1.83E-11	AAEL010159-PA OS=Aedes aegypti GN=AAEL010159 PE=4 SV=1
CCG013071.1	1207	3507	26595	124.850753	770.2695698	2.625158905	Up	1.79E-11	3.93E-11	Putative salivary secreted peptide of the 13 kDa family OS=Aedes triseriatus PE=2 SV=1
CCG008270.1	577	6949	26623	517.498589	1612.988219	1.640109068	Up	1.69E-12	4.05E-12	Ubiquitin OS=Anopheles stephensi GN=Ubi-p63E PE=2 SV=1
CCG027422.1	940	8370	26964	382.613488	1002.781888	1.390048206	Up	0	0	AAEL000269-PA OS=Aedes aegypti GN=AAEL000269 PE=4 SV=1
CCG023199.2	13053	2064	27293	6.79457077	73.09555259	3.42732931	Up	0	0	AGAP000029-PA OS=Anopheles gambiae GN=AgaP_AGAP000029 PE=4 SV=5
CCG022904.1	9326	9917	28144	45.6927791	105.4970771	1.207164931	Up	0	0	Mitochondrial phosphate carrier protein OS=Aedes aegypti PE=2 SV=1
CCG007010.1	784	58379	28439	3199.65579	1268.084736	-1.335265558	Down	0	0	Putative uncharacterized protein OS=Daphnia pulex GN=DAPPUDRAFT_67491 PE=4 SV=1
CCG011189.1	9303	4375	28942	20.2077382	108.7565781	2.428122903	Up	3.03E-11	6.59E-11	AAEL015015-PA OS=Aedes aegypti GN=AAEL015015 PE=4 SV=1

CCG003364.1	2292	0	29097	0.001	443.7962388	18.75953791	Up	4.44E-16	1.66E-15	AAEL004401-PA (Fragment) OS=Aedes aegypti GN=HPX7 PE=4 SV=2
CCG002555.1	4217	3819	29479	38.9142557	244.3763436	2.650733964	Up	6.38E-12	1.44E-11	AAEL007880-PA OS=Aedes aegypti GN=AAEL007880 PE=3 SV=1
CCG006336.1	1002	7714	30004	330.806919	1046.794547	1.661917	Up	2.48E-11	5.41E-11	AAEL007699-PA OS=Aedes aegypti GN=AAEL007699 PE=2 SV=1
CCG016778.1	1470	4844	30352	141.595506	721.8051814	2.349834018	Up	1.38E-11	3.05E-11	AAEL009994-PA OS=Aedes aegypti GN=AAEL009994 PE=2 SV=1
CCG002408.1	1447	7797	30417	231.537678	734.8485864	1.666200033	Up	0	0	Ribosomal protein S5 OS=Aedes albopictus PE=2 SV=1
CCG015165.1	6081	9995	30436	70.6269524	174.9695958	1.308813499	Up	2.77E-11	6.04E-11	AAEL004716-PA OS=Aedes aegypti GN=AAEL004716 PE=4 SV=1
CCG026845.1	7798	10253	30633	56.4976518	137.3271214	1.281353766	Up	0	0	AAEL005515-PA OS=Aedes aegypti GN=AAEL005515 PE=4 SV=1
CCG008943.1	2830	5878	31114	89.2495053	384.3434005	2.106479819	Up	2.52E-11	5.51E-11	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003009 PE=4 SV=1
CCG002644.1	822	5581	31567	291.744634	1342.49139	2.202134812	Up	6.47E-12	1.46E-11	Ribosomal protein S2 OS=Aedes albopictus PE=2 SV=1
CCG022918.1	5323	4178	31623	33.7267615	207.6809292	2.622403046	Up	1.05E-11	2.33E-11	AAEL002551-PA OS=Aedes aegypti GN=AAEL002551 PE=4 SV=1
CCG004868.1	5466	0	32075	0.001	205.1384426	17.64623835	Up	4.44E-16	1.65E-15	AAEL007657-PA OS=Aedes aegypti GN=AAEL007657 PE=4 SV=1
CCG011071.1	5103	3999	32157	33.6735192	220.292633	2.709734838	Up	5.80E-12	1.31E-11	AAEL001898-PA OS=Aedes aegypti GN=AAEL001898 PE=4 SV=1
CCG014955.1	2342	6158	32532	112.983615	485.5946456	2.10363895	Up	0	0	Nucleolar GTP-binding protein 1 OS=Aedes aegypti GN=AAEL009859 PE=3 SV=1
CCG005172.1	3803	10575	33093	119.485917	304.2004347	1.348181624	Up	0	0	Heat shock cognate 70 OS=Aedes aegypti PE=2 SV=1
CCG000428.4	3609	11865	33513	141.267914	324.6208727	1.200321942	Up	0	0	AAEL000804-PA OS=Aedes aegypti GN=AAEL000804 PE=4 SV=1
CCG008274.1	2592	142	33738	2.35405182	455.0240516	7.594652917	Up	2.62E-13	7.34E-13	AAEL002422-PA OS=Aedes aegypti GN=AAEL002422 PE=4 SV=1

CCG028312.1	4062	9772	34847	103.372784	299.8993281	1.536621894	Up	2.40E-11	5.23E-11	AAEL011104-PA OS=Aedes aegypti GN=AAEL011104 PE=4 SV=1
CCG007988.1	46271	7610	36202	7.06705454	27.35103057	1.952414247	Up	0	0	AAEL006959-PA OS=Aedes aegypti GN=AAEL006959 PE=4 SV=1
CCG003152.1	2105	8872	37379	181.105694	620.7626734	1.777209906	Up	0	0	Heat shock cognate 70 OS=Aedes aegypti PE=2 SV=1
CCG012273.2	3687	141	38538	1.64326894	365.3979035	7.796755933	Up	0	0	AAEL004388-PA (Fragment) OS=Aedes aegypti GN=HPX8A PE=4 SV=2
CCG008581.1	638	13230	38916	891.049512	2132.345296	1.258863572	Up	0	0	AAEL007699-PA OS=Aedes aegypti GN=AAEL007699 PE=2 SV=1
CCG007256.1	1305	7976	39400	262.62575	1055.445291	2.006771519	Up	0	0	Ribosomal protein L3 OS=Aedes aegypti PE=2 SV=1
CCG007978.1	2841	783	42786	11.8427674	526.4782928	5.474295896	Up	0	0	AAEL004978-PA OS=Aedes aegypti GN=AAEL004978 PE=2 SV=1
CCG021835.1	6417	16676	43134	111.666401	234.9836856	1.073365436	Up	0	0	AAEL017349-PA OS=Aedes aegypti PE=2 SV=1
CCG007823.1	429	63	44252	6.31024075	3605.999216	9.15848832	Up	0	0	Chymotrypsin-like serine protease OS=Aedes aegypti GN=JA15 PE=2 SV=1
CCG019426.1	405	12826	44651	1360.81436	3854.128435	1.501934378	Up	0	0	OS=ribosomal protein L32 OS=Culex quinquefasciatus GN=CpipJ_CPIJ001220 PE=4 SV=1
CCG002645.1	6483	8832	45858	58.5390554	247.2800642	2.078674563	Up	0	0	AAEL010159-PA OS=Aedes aegypti GN=AAEL010159 PE=4 SV=1
CCG010291.1	1320	3185	46519	103.680761	1231.987937	3.57076801	Up	0	0	AAEL008500-PA OS=Aedes aegypti GN=AAEL008500 PE=3 SV=1
CCG026423.1	9565	2329	48385	10.4627822	176.8380786	4.079090521	Up	0	0	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009795 PE=4 SV=1
CCG015184.1	1699	33	48546	0.83460932	998.8725746	10.22498389	Up	1.31E-13	4.01E-13	AAEL000442-PA OS=Aedes aegypti GN=AAEL000442 PE=4 SV=1
CCG012092.1	1809	249	50992	5.91457375	985.4021278	7.380294576	Up	0	0	AAEL010097-PA OS=Aedes aegypti GN=AAEL010097 PE=4 SV=1

CCG001066.1	2088	320333	52037	6592.25287	871.2278967	-2.919649529	Down	0	0	ATP synthase subunit alpha OS=Aedes aegypti GN=AAEL012175 PE=2 SV=1
CCG026558.1	8283	401328	55728	2081.97002	235.1991848	-3.145994327	Down	0	0	AAEL005733-PA OS=Aedes aegypti GN=AAEL005733 PE=4 SV=1
CCG027751.1	1795	16975	56883	406.35724	1107.817079	1.446899184	Up	0	0	Elongation factor 1-gamma OS=Aedes aegypti PE=2 SV=1
CCG014051.1	661	17515	58235	1138.60046	3079.871782	1.435608708	Up	0	0	60S ribosomal protein L17 OS=Culex quinquefasciatus GN=CpipJ_CPIJ015624 PE=3 SV=1
CCG024781.1	985	25766	73431	1124.01846	2606.112915	1.213233865	Up	1.43E-10	3.04E-10	60S ribosomal protein L31 OS=Culex quinquefasciatus GN=CpipJ_CPIJ010692 PE=3 SV=1
CCG011426.1	813	155	74433	8.19226183	3200.552786	8.609843563	Up	3.42E-13	9.26E-13	AAEL003060-PA (Precursor) OS=Aedes aegypti GN=CHYM PE=2 SV=1
CCG016979.2	2286	24382	76430	458.306242	1168.793129	1.350635762	Up	4.23E-11	9.13E-11	AAEL010787-PA OS=Aedes aegypti GN=AAEL010787 PE=3 SV=1
CCG003471.1	1389	31475	78256	973.70223	1969.54277	1.016308196	Up	1.70E-10	3.62E-10	AAEL008188-PA OS=Aedes aegypti GN=AAEL008188 PE=4 SV=1
CCG015935.1	813	20	89892	1.05706604	3865.276034	11.83629022	Up	3.69E-10	7.70E-10	AAEL003060-PA (Precursor) OS=Aedes aegypti GN=CHYM PE=2 SV=1
CCG016661.1	51084	238399	91016	200.531316	62.28491345	-1.686872892	Down	0	0	AAEL006582-PA OS=Aedes aegypti GN=AAEL006582 PE=3 SV=1
CCG012274.1	2355	21	91549	0.38316961	1358.978888	11.79225229	Up	1.76E-10	3.73E-10	AAEL004388-PA (Fragment) OS=Aedes aegypti GN=HPX8A PE=4 SV=2
CCG014531.2	948	26855	102590	1217.24918	3783.09031	1.635940688	Up	1.61E-10	3.43E-10	AAEL010821-PA OS=Aedes aegypti GN=AAEL010821 PE=2 SV=1
CCG014436.2	2494	36874	133320	635.311144	1868.73996	1.556530596	Up	2.25E-10	4.75E-10	AGAP006023-PA OS=Anopheles gambiae GN=AGAP006023 PE=4 SV=3
CCG017696.1	1459	31481	138098	927.162588	3308.888352	1.835452356	Up	2.02E-10	4.26E-10	AAEL006259-PA OS=Aedes aegypti GN=GPROP2 PE=3 SV=1

CCG003493.1	696	51	144493	3.14864435	7257.509031	11.17052794	Up	0	0	Trypsin OS=Ochlerotatus taeniorhynchus PE=2 SV=2
CCG025915.1	759	40	210225	2.26454464	9682.613616	12.06195983	Up	6.86E-14	2.24E-13	Trypsin OS=Ochlerotatus taeniorhynchus PE=2 SV=2