

Table S7.3. Candidate female-specific 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
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
CCG015183.1	309	0	2985	0.001	337.7037416	18.36539864	Up	4.44E-16	1.65E-15	-
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
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
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
CCG021499.1	633	0	3183	0.001	175.7854581	17.4234562	Up	4.44E-16	1.63E-15	30 kDa salivary gland allergen 30k-2 OS=Aedes albopictus PE=2 SV=1
CCG027373.1	761	0	3413	0.001	156.7839667	17.25841851	Up	4.44E-16	1.64E-15	-
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
CCG007095.1	264	0	663	0.001	87.79294508	16.42181739	Up	4.44E-16	1.65E-15	-
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
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
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
CCG015620.1	744	0	1246	0.001	58.54571621	15.83727599	Up	4.44E-16	1.65E-15	AAEL007855-PA OS=Aedes aegypti GN=AAEL007855 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
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
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

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
CCG015513.1	542	0	837	0.001	53.98538313	15.72028122	Up	4.44E-16	1.64E-15	-
CCG013230.1	481	0	698	0.001	50.7294744	15.63053659	Up	4.44E-16	1.63E-15	-
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
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
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
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
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
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
CCG020758.1	738	0	817	0.001	38.70042293	15.24006171	Up	4.44E-16	1.65E-15	Serine 3-dehydrogenase OS=Culex quinquefasciatus GN=CpipJ_CPIJ016922 PE=3 SV=1
CCG006000.1	641	0	669	0.001	36.48531538	15.1550283	Up	4.44E-16	1.65E-15	-
CCG027682.1	579	0	556	0.001	33.56960602	15.03486798	Up	4.44E-16	1.65E-15	AAEL008835-PA OS=Aedes aegypti GN=AAEL008835 PE=4 SV=1
CCG025737.1	441	0	422	0.001	33.45213786	15.02981079	Up	4.44E-16	1.64E-15	-
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
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
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
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
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
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
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

CCG001793.1	363	0	271	0.001	26.09832784	14.67166975	Up	4.44E-16	1.64E-15	-
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
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
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
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
CCG004157.1	354	0	255	0.001	25.18180954	14.62009434	Up	4.44E-16	1.63E-15	-
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
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
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
CCG005597.1	261	0	170	0.001	22.76975882	14.47483139	Up	4.44E-16	1.66E-15	-
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
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
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
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
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
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
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
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
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

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
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
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
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
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
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
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
CCG018501.1	1558	0	698	0.001	15.661667	13.93495016	Up	4.44E-16	1.64E-15	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 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
CCG026935.1	444	0	195	0.001	15.35329723	13.9062609	Up	4.44E-16	1.63E-15	AAEL003185-PA OS=Aedes aegypti GN=AAEL003185 PE=4 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
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
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
CCG003214.1	594	0	233	0.001	13.71259005	13.74321347	Up	4.44E-16	1.63E-15	AAEL006892-PA OS=Aedes aegypti GN=AAEL006892 PE=4 SV=1
CCG008458.1	513	0	190	0.001	12.94750992	13.66038704	Up	4.44E-16	1.64E-15	AAEL001949-PA OS=Aedes aegypti GN=AAEL001949 PE=4 SV=1
CCG025837.1	273	0	101	0.001	12.93328188	13.65880079	Up	4.44E-16	1.65E-15	-
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
CCG022213.1	594	0	195	0.001	11.47620197	13.48635764	Up	4.44E-16	1.66E-15	AAEL006892-PA OS=Aedes aegypti GN=AAEL006892 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
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
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
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
CCG007503.1	730	0	213	0.001	10.20015473	13.31630342	Up	4.44E-16	1.64E-15	AAEL012453-PA OS=Aedes aegypti GN=AAEL012453 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
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
CCG007454.1	318	0	84	0.001	9.234261789	13.17278092	Up	4.44E-16	1.65E-15	-
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
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
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
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
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
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
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
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
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
CCG011355.1	621	0	136	0.001	7.655918907	12.90235983	Up	4.44E-16	1.65E-15	-
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

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
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
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
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
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
CCG005992.1	636	0	122	0.001	6.705832966	12.71120083	Up	4.44E-16	1.65E-15	-
CCG021098.1	491	0	94	0.001	6.692623252	12.70835609	Up	4.44E-16	1.64E-15	-
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
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
CCG015867.1	1667	0	306	0.001	6.417056204	12.6476959	Up	4.44E-16	1.64E-15	AAEL006123-PA OS=Aedes aegypti GN=AAEL006123 PE=3 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
CCG001796.1	363	0	63	0.001	6.067138944	12.56680064	Up	4.44E-16	1.65E-15	-
CCG014303.1	708	0	120	0.001	5.925131656	12.5326315	Up	4.44E-16	1.63E-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
CCG018919.1	969	2	13919	0.0886888	502.1509333	12.46708115	Up	9.36E-05	0.000153	AAEL000923-PA OS=Aedes aegypti GN=AAEL000923 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
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
CCG008065.1	351	0	55	0.001	5.477792657	12.41937894	Up	6.66E-15	2.40E-14	-
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
CCG007674.1	519	0	77	0.001	5.186488076	12.34054226	Up	4.44E-16	1.64E-15	-
CCG009456.1	513	0	76	0.001	5.179003966	12.33845895	Up	4.44E-16	1.63E-15	-

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
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
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
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
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
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
CCG007675.1	648	0	80	0.001	4.315836639	12.07542454	Up	4.44E-16	1.65E-15	AAEL011896-PA OS=Aedes aegypti GN=AAEL011896 PE=4 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
CCG007694.1	304	0	37	0.001	4.254790265	12.0548723	Up	3.00E-10	6.30E-10	Importin9 OS=Culex quinquefasciatus GN=CpipJ_CPIJ008197 PE=4 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
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
CCG017704.1	307	0	35	0.001	3.985471293	11.96053462	Up	9.86E-10	2.03E-09	-
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
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
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
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
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
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

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
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
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
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
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
CCG003434.1	764	0	81	0.001	3.706309448	11.85576763	Up	4.44E-16	1.64E-15	-
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
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
CCG003332.1	1017	0	105	0.001	3.609261614	11.817488	Up	4.44E-16	1.63E-15	AAEL009198-PB OS=Aedes aegypti GN=AAEL009198 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
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
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
CCG015454.1	852	0	85	0.001	3.487621509	11.76802777	Up	4.44E-16	1.63E-15	AAEL003128-PA OS=Aedes aegypti GN=AAEL003128 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
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
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
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
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

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
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
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
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
CCG020332.1	273	0	23	0.001	2.945202805	11.52415127	Up	1.25E-06	2.29E-06	-
CCG025524.1	672	0	56	0.001	2.913189731	11.50838395	Up	3.77E-15	1.36E-14	-
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
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
CCG003930.1	354	0	28	0.001	2.76506144	11.43309582	Up	6.36E-08	1.23E-07	-
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
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
CCG016419.1	689	0	52	0.001	2.638360511	11.365426	Up	3.95E-14	1.36E-13	-
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
CCG004395.1	228	0	17	0.001	2.60653818	11.34791928	Up	4.44E-05	7.47E-05	-
CCG000318.1	297	0	22	0.001	2.589501983	11.33845895	Up	2.26E-06	4.09E-06	-
CCG014440.1	435	0	32	0.001	2.571643349	11.32847486	Up	5.88E-09	1.18E-08	-
CCG003756.1	390	0	28	0.001	2.509824999	11.29337106	Up	6.36E-08	1.23E-07	-
CCG016848.1	1173	0	84	0.001	2.503406009	11.28967657	Up	4.44E-16	1.63E-15	AAEL008858-PA OS=Aedes aegypti GN=AAEL008858 PE=4 SV=1
CCG007744.1	1461	0	102	0.001	2.440618912	11.25303133	Up	4.44E-16	1.63E-15	AAEL006058-PA OS=Aedes aegypti GN=CYP325Q2 PE=3 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
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
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
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
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
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
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
CCG011427.1	576	0	36	0.001	2.184892298	11.09334645	Up	5.44E-10	1.13E-09	-
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
CCG003581.1	1773	0	109	0.001	2.149155199	11.06955395	Up	4.44E-16	1.65E-15	-
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
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
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
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
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
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
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
CCG023894.1	768	0	44	0.001	2.00281794	10.96781557	Up	4.65E-12	1.06E-11	-
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
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
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
CCG018745.1	291	0	16	0.001	1.922104565	10.90847111	Up	8.05E-05	0.000133	AAEL001787-PA OS=Aedes aegypti GN=AAEL001787 PE=4 SV=1

CCG011485.1	787	0	43	0.001	1.910045618	10.89939138	Up	8.43E-12	1.88E-11	-
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
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
CCG002213.1	1042	0	56	0.001	1.878755757	10.87556181	Up	3.77E-15	1.36E-14	-
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
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
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
CCG006420.1	1434	1	2280	0.0299649	55.58219738	10.85713131	Up	0.00059	0.000916	Tubulin alpha-2 chain OS=Culex quinquefasciatus GN=CpipJ_CPIJ004534 PE=3 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
CCG020646.1	271	0	14	0.001	1.805962638	10.81855233	Up	0.00026	0.000423	-
CCG005210.1	935	1	2216	0.0459569	82.85298538	10.81605537	Up	0.00059	0.000918	AAEL017320-PA OS=Aedes aegypti GN=FOHSDR-2 PE=2 SV=1
CCG019640.1	315	0	16	0.001	1.775658503	10.79413843	Up	8.05E-05	0.000133	AAEL006670-PA OS=Aedes aegypti GN=AAEL006670 PE=4 SV=1
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
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
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
CCG015948.1	282	0	14	0.001	1.735517287	10.76115002	Up	0.00026	0.000423	AAEL005702-PA OS=Aedes aegypti GN=AAEL005702 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

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
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
CCG028030.1	531	0	25	0.001	1.645869905	10.68463459	Up	3.80E-07	7.11E-07	-
CCG006085.3	1983	1	2001	0.0216691	35.2755985	10.66881866	Up	0.00059	0.000917	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1
CCG020221.1	345	0	16	0.001	1.621253416	10.6628939	Up	8.05E-05	0.000133	-
CCG002901.1	1123	0	52	0.001	1.618726974	10.66064396	Up	3.95E-14	1.36E-13	-
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
CCG023240.1	309	0	14	0.001	1.583870145	10.62923834	Up	0.00026	0.000423	AAEL012826-PA OS=Aedes aegypti GN=AAEL012826 PE=4 SV=1
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
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
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
CCG009949.1	381	0	16	0.001	1.468064116	10.51969926	Up	8.05E-05	0.000133	AAEL000073-PA OS=Aedes aegypti GN=AAEL000073 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
CCG027450.1	669	0	28	0.001	1.463126681	10.51483897	Up	6.36E-08	1.23E-07	-
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
CCG004491.1	362	0	15	0.001	1.44854738	10.50039116	Up	0.00015	0.000236	-
CCG023219.1	388	0	16	0.001	1.441578424	10.49343361	Up	8.05E-05	0.000133	AAEL010576-PM OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG006370.1	853	0	35	0.001	1.434395882	10.48622754	Up	9.86E-10	2.03E-09	-
CCG001247.1	390	0	16	0.001	1.434185714	10.48601614	Up	8.05E-05	0.000133	-
CCG001883.1	342	0	14	0.001	1.43104057	10.48284886	Up	0.00026	0.000423	-
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

CCG006392.1	1683	2	3430	0.0510633	71.24592355	10.44630606	Up	9.36E-05	0.000154	AAEL000335-PA OS=Aedes aegypti GN=AAEL000335 PE=4 SV=1
CCG027975.1	351	0	14	0.001	1.394347222	10.44537415	Up	0.00026	0.000423	-
CCG007233.1	429	0	17	0.001	1.385293019	10.43597545	Up	4.44E-05	7.48E-05	-
CCG014996.1	357	0	14	0.001	1.370912815	10.42092111	Up	0.00026	0.000423	-
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
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
CCG003228.1	573	0	22	0.001	1.342202599	10.39038674	Up	2.26E-06	4.09E-06	Modifier 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
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
CCG025882.1	348	0	13	0.001	1.305912638	10.35084267	Up	0.00048	0.000755	AAEL013443-PA OS=Aedes aegypti GN=AAEL013443 PE=4 SV=1
CCG015725.1	1644	1	1593	0.0261373	33.87380468	10.33984376	Up	0.00059	0.000918	AAEL000889-PA (Fragment) OS=Aedes aegypti GN=AAEL000889 PE=4 SV=2
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 / SRRRC 167) GN=AFLA_063980 PE=4 SV=1
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
CCG015990.1	741	0	27	0.001	1.273783364	10.31490422	Up	1.15E-07	2.21E-07	Trypsin-2 OS=Culex quinquefasciatus GN=CpipJ_CPIJ000771 PE=3 SV=1
CCG006089.1	1143	1	1531	0.0375938	46.82512838	10.28257177	Up	0.00059	0.000916	AAEL005938-PA OS=Aedes aegypti GN=AAEL005938 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
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
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

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
CCG004140.1	528	0	18	0.001	1.191759435	10.21887733	Up	2.45E-05	4.20E-05	-
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
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
CCG028048.1	848	0	28	0.001	1.154282724	10.17278092	Up	6.36E-08	1.23E-07	-
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
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
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
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
CCG014442.1	507	0	16	0.001	1.10321978	10.10750451	Up	8.05E-05	0.000133	AGAP004106-PA OS=Anopheles gambiae GN=AgaP_AGAP004106 PE=4 SV=5
CCG018521.1	480	0	15	0.001	1.092446149	10.09334645	Up	0.00015	0.000236	AGAP013413-PA OS=Anopheles gambiae GN=AgaP_AGAP013413 PE=4 SV=1
CCG018342.1	513	0	16	0.001	1.090316624	10.09053143	Up	8.05E-05	0.000133	-
CCG011352.1	905	2	2656	0.0949607	102.5957824	10.07735264	Up	9.36E-05	0.000154	Cation efflux protein/ zinc transporter OS=Culex quinquefasciatus GN=CpipJ_CPIJ010101 PE=4 SV=1
CCG007531.1	1347	2	2622	0.0638006	68.04795969	10.05876517	Up	9.36E-05	0.000154	AAEL004172-PA OS=Aedes aegypti GN=AAEL004172 PE=3 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
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
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
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
CCG014603.1	504	0	15	0.001	1.040424904	10.02295712	Up	0.00015	0.000236	AAEL011252-PA OS=Aedes aegypti GN=AAEL011252 PE=4 SV=1

CCG009113.1	445	0	13	0.001	1.021253029	9.996124642	Up	0.00048	0.000756	-
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
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
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
CCG001329.1	462	0	13	0.001	0.983674455	9.942037127	Up	0.00048	0.000756	AAEL004611-PA OS=Aedes aegypti GN=AAEL004611 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
CCG003908.1	714	0	20	0.001	0.979223439	9.935494281	Up	7.45E-06	1.32E-05	-
CCG017910.1	465	0	13	0.001	0.977328168	9.932699262	Up	0.00048	0.000755	AAEL004145-PA OS=Aedes aegypti GN=AAEL004145 PE=4 SV=1
CCG002251.1	576	0	16	0.001	0.971063244	9.923421449	Up	8.05E-05	0.000133	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 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
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
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
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
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
CCG010936.1	486	0	13	0.001	0.935097938	9.868973665	Up	0.00048	0.000756	-
CCG012873.1	834	1	1149	0.0515225	48.16194246	9.868476286	Up	0.00059	0.000917	AAEL014128-PA OS=Aedes aegypti GN=AAEL014128 PE=4 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
CCG028611.1	528	0	14	0.001	0.926924005	9.856307253	Up	0.00026	0.000423	AAEL011712-PA OS=Aedes aegypti GN=AAEL011712 PE=4 SV=1
CCG023853.1	1434	0	38	0.001	0.926369956	9.855444655	Up	1.65E-10	3.51E-10	AAEL008538-PA OS=Aedes aegypti GN=AAEL008538 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
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
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
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
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
CCG008688.1	507	0	13	0.001	0.896366071	9.807944231	Up	0.00048	0.000756	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 SV=1
CCG009755.1	664	0	17	0.001	0.895016122	9.80576986	Up	4.44E-05	7.47E-05	-
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
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
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
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
CCG012832.1	648	0	16	0.001	0.863167328	9.753496447	Up	8.05E-05	0.000133	AAEL003547-PA OS=Aedes aegypti GN=AAEL003547 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
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
CCG002798.1	538	0	13	0.001	0.844716725	9.722323806	Up	0.00048	0.000755	-
CCG021779.1	668	0	16	0.001	0.837323995	9.709642158	Up	8.05E-05	0.000133	-
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
CCG022897.2	688	0	16	0.001	0.812983181	9.667081695	Up	8.05E-05	0.000133	-
CCG026145.1	649	2	1995	0.1324183	107.4603423	9.664486235	Up	9.36E-05	0.000154	AAEL009504-PA OS=Aedes aegypti GN=AAEL009504 PE=4 SV=1
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

CCG028412.1	696	0	16	0.001	0.803638546	9.650402954	Up	8.05E-05	0.000133	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ000496 PE=4 SV=1
CCG021703.1	654	0	15	0.001	0.801795339	9.64709022	Up	0.00015	0.000236	AAEL010547-PA OS=Aedes aegypti GN=AAEL010547 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
CCG025315.1	905	0	20	0.001	0.772558603	9.593500563	Up	7.45E-06	1.32E-05	-
CCG009511.1	600	0	13	0.001	0.75742933	9.564967478	Up	0.00048	0.000755	AAEL001383-PA OS=Aedes aegypti GN=AAEL001383 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
CCG022386.1	604	0	13	0.001	0.752413242	9.555381429	Up	0.00048	0.000755	-
CCG007668.1	1260	0	27	0.001	0.749105931	9.549025934	Up	1.15E-07	2.21E-07	-
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
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
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
CCG021545.1	1639	2	1818	0.0524341	38.77617277	9.530449688	Up	9.36E-05	0.000154	AAEL002638-PA OS=Aedes aegypti GN=CYP9J6 PE=3 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
CCG004271.1	670	0	14	0.001	0.730471455	9.512684087	Up	0.00026	0.000423	Gag polyprotein OS=Anopheles gambiae 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
CCG010484.1	897	1	876	0.0479038	34.13985558	9.477100263	Up	0.00059	0.000916	AAEL013295-PA OS=Aedes aegypti GN=AAEL013295 PE=4 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
CCG018585.3	1455	0	29	0.001	0.696762905	9.444524007	Up	3.51E-08	6.85E-08	AAEL005963-PA OS=Aedes aegypti GN=CASPS15 PE=3 SV=1
CCG023512.1	756	0	15	0.001	0.693616603	9.437994622	Up	0.00015	0.000236	AAEL005465-PA OS=Aedes aegypti GN=AAEL005465 PE=4 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

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
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
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
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
CCG009507.1	732	0	14	0.001	0.668600922	9.385001534	Up	0.00026	0.000423	AAEL014647-PA OS=Aedes aegypti GN=AAEL014647 PE=4 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
CCG023752.1	1080	2	1631	0.0795736	52.79347168	9.373854271	Up	9.36E-05	0.000153	AAEL009263-PA OS=Aedes aegypti GN=AAEL009263 PE=2 SV=1
CCG019765.1	693	0	13	0.001	0.65578297	9.357074626	Up	0.00048	0.000755	AAEL004751-PA OS=Aedes aegypti GN=AAEL004751 PE=4 SV=1
CCG008302.1	1098	2	1605	0.0782691	51.10021331	9.350670786	Up	9.36E-05	0.000154	AAEL010638-PA OS=Aedes aegypti GN=AAEL010638 PE=3 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
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
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
CCG011354.1	551	1	780	0.077985	49.48721576	9.309643518	Up	0.00059	0.000918	-
CCG028209.1	776	0	14	0.001	0.63069056	9.30078853	Up	0.00026	0.000423	AAEL009165-PA OS=Aedes aegypti GN=AAEL009165 PE=4 SV=1
CCG028277.1	834	0	15	0.001	0.628745985	9.296333472	Up	0.00015	0.000237	AAEL006378-PA OS=Aedes aegypti GN=AAEL006378 PE=3 SV=1
CCG000163.1	788	0	14	0.001	0.621086136	9.278649553	Up	0.00026	0.000423	AAEL009165-PA OS=Aedes aegypti GN=AAEL009165 PE=4 SV=1
CCG000733.1	794	0	14	0.001	0.616392789	9.267706175	Up	0.00026	0.000423	AAEL013652-PA OS=Aedes aegypti GN=AAEL013652 PE=3 SV=1
CCG009150.1	744	0	13	0.001	0.610830105	9.254627357	Up	0.00048	0.000755	-

CCG018105.1	900	2	1487	0.0954883	57.75884173	9.240502136	Up	9.36E-05	0.000153	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ015628 PE=4 SV=1
CCG021535.1	1296	2	1476	0.0663113	39.81359299	9.22979021	Up	9.36E-05	0.000154	AAEL002586-PA OS=Aedes aegypti GN=AAEL002586 PE=4 SV=1
CCG028103.1	816	0	14	0.001	0.599774356	9.22827603	Up	0.00026	0.000423	AAEL005607-PA OS=Aedes aegypti GN=AAEL005607 PE=3 SV=1
CCG012066.1	825	0	14	0.001	0.593231363	9.212451063	Up	0.00026	0.000423	AAEL012564-PA OS=Aedes aegypti GN=AAEL012564 PE=3 SV=1
CCG004721.2	832	0	14	0.001	0.588240234	9.200261654	Up	0.00026	0.000423	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014613 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
CCG022299.1	994	0	16	0.001	0.56270868	9.136244409	Up	8.05E-05	0.000133	Putative uncharacterized protein (Fragment) OS=Harpegnathos saltator GN=EAI_06049 PE=4 SV=1
CCG010650.1	446	1	689	0.0963447	54.00505089	9.130673377	Up	0.00059	0.000917	Chromosome undetermined SCAF14565, whole genome shotgun sequence OS=Tetraodon nigroviridis GN=GSTENG00016696001 PE=3 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
CCG004308.1	951	0	15	0.001	0.551392378	9.106935515	Up	0.00015	0.000237	AAEL010140-PA OS=Aedes aegypti GN=AAEL010140 PE=3 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
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
CCG013061.1	588	2	1305	0.1461556	77.58597141	9.052147295	Up	9.36E-05	0.000153	AAEL008182-PA OS=Aedes aegypti GN=AAEL008182 PE=4 SV=1
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
CCG002815.1	1216	0	18	0.001	0.517474492	9.015343938	Up	2.45E-05	4.20E-05	-
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
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	FAD oxidoreductase OS=Culex quinquefasciatus GN=CpipJ_CPIJ016820 PE=4 SV=1
CCG022378.1	1116	0	16	0.001	0.501193932	8.969225138	Up	8.05E-05	0.000133	AAEL008380-PA OS=Aedes aegypti GN=AAEL008380 PE=4 SV=1
CCG022315.1	981	0	14	0.001	0.498894877	8.962592046	Up	0.00026	0.000423	Uncharacterized protein OS=Anopheles darlingi GN=AND_21982 PE=4 SV=1
CCG000478.1	1134	0	16	0.001	0.493238473	8.946141525	Up	8.05E-05	0.000133	AAEL000130-PA OS=Aedes aegypti GN=AAEL000130 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
CCG022990.1	1267	0	17	0.001	0.469053437	8.873608482	Up	4.44E-05	7.47E-05	Endonuclease-reverse transcriptase OS=Bombyx mori PE=4 SV=1
CCG025268.1	972	0	13	0.001	0.467548969	8.868973665	Up	0.00048	0.000756	AAEL007999-PA OS=Aedes aegypti GN=AAEL007999 PE=4 SV=1
CCG021767.1	1050	0	14	0.001	0.466110357	8.86452776	Up	0.00026	0.000423	Putative uncharacterized protein (Fragment) OS=Camponotus floridanus GN=EAG_00896 PE=4 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
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
CCG020738.1	846	1	562	0.0507916	23.22287417	8.836739524	Up	0.00059	0.000918	AAEL010318-PA OS=Aedes aegypti GN=AAEL010318 PE=2 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
CCG024753.1	981	1	558	0.043802	19.8845244	8.826434516	Up	0.00059	0.000918	AAEL006884-PA OS=Aedes aegypti GN=AAEL006884 PE=4 SV=1
CCG010649.1	464	1	553	0.0926072	41.66363589	8.813448874	Up	0.00059	0.000916	Histone H2A OS=Gasterosteus aculeatus PE=3 SV=1
CCG003766.1	1254	0	16	0.001	0.446038619	8.801024817	Up	8.05E-05	0.000133	AAEL001160-PA OS=Aedes aegypti GN=AAEL001160 PE=4 SV=1
CCG011758.1	1188	0	15	0.001	0.441392383	8.785917925	Up	0.00015	0.000236	Odorant receptor 2 OS=Aedes albopictus PE=2 SV=1
CCG003137.1	1032	0	13	0.001	0.44036589	8.782558913	Up	0.00048	0.000756	AAEL015565-PA OS=Aedes aegypti GN=AAEL015565 PE=4 SV=1
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
CCG005674.1	1344	0	16	0.001	0.416169962	8.701029027	Up	8.05E-05	0.000133	-

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
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
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
CCG021419.1	1152	0	13	0.001	0.394494443	8.623861167	Up	0.00048	0.000756	AAEL010973-PA OS=Aedes aegypti GN=AAEL010973 PE=4 SV=1
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
CCG003429.1	1257	0	14	0.001	0.389352327	8.604932438	Up	0.00026	0.000423	AAEL009473-PA OS=Aedes aegypti GN=AAEL009473 PE=4 SV=1
CCG019144.1	1170	0	13	0.001	0.388425297	8.601493354	Up	0.00048	0.000756	AAEL006371-PA OS=Aedes aegypti GN=AAEL006371 PE=3 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
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
CCG026474.1	362	2	874	0.2374018	84.40202735	8.473802673	Up	9.36E-05	0.000153	Histone H2A OS=Drosophila willistoni GN=GK10379 PE=3 SV=1
CCG021055.1	1398	0	14	0.001	0.350082886	8.451552727	Up	0.00026	0.000423	Putative uncharacterized protein (Fragment) OS=Solenopsis invicta GN=SINV_08164 PE=4 SV=1
CCG014381.1	1608	0	16	0.001	0.34784355	8.442294759	Up	8.05E-05	0.000133	AAEL006875-PA OS=Aedes aegypti GN=CYP307B1_a PE=3 SV=1
CCG011353.1	736	2	849	0.1167656	40.3255122	8.431933947	Up	9.36E-05	0.000154	-
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
CCG000380.1	1176	2	808	0.0730778	24.01895207	8.360524687	Up	9.36E-05	0.000153	AAEL005606-PA OS=Aedes aegypti GN=AAEL005606 PE=4 SV=1
CCG027428.1	1491	0	14	0.001	0.32824673	8.35863683	Up	0.00026	0.000423	AAEL005151-PA OS=Aedes aegypti GN=AAEL005151 PE=4 SV=1
CCG024624.1	398	1	402	0.1079642	35.30961624	8.353364895	Up	0.00059	0.000915	Chromosome undetermined SCAF14565, whole genome shotgun sequence OS=Tetraodon nigroviridis GN=GSTENG00016696001 PE=3 SV=1
CCG018391.1	777	2	800	0.1106042	35.99307776	8.346169394	Up	9.36E-05	0.000153	-

CCG023244.1	1533	0	14	0.001	0.319253669	8.318559391	Up	0.00026	0.000423	AAEL007377-PA OS=Aedes aegypti GN=AAEL007377 PE=4 SV=1
CCG016125.1	1170	2	775	0.0734525	23.1561235	8.300365704	Up	9.36E-05	0.000154	AAEL006726-PA OS=Aedes aegypti GN=AAEL006726 PE=4 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
CCG009856.1	1179	2	760	0.0728918	22.53459741	8.272168812	Up	9.36E-05	0.000154	AAEL012453-PA OS=Aedes aegypti GN=AAEL012453 PE=4 SV=1
CCG019130.1	734	1	378	0.0585419	18.00303627	8.264555628	Up	0.00059	0.000916	AAEL009973-PA OS=Aedes aegypti GN=AAEL009973 PE=4 SV=1
CCG016477.1	489	1	377	0.0878727	26.95147309	8.260733917	Up	0.00059	0.000916	AAEL013262-PA OS=Aedes aegypti GN=AAEL013262 PE=4 SV=1
CCG025364.1	1593	0	13	0.001	0.285284117	8.156255617	Up	0.00048	0.000755	AAEL009243-PA OS=Aedes aegypti GN=AAEL009243 PE=3 SV=1
CCG017133.1	1329	2	699	0.0646648	18.38663316	8.151461849	Up	9.36E-05	0.000153	AAEL009987-PA OS=Aedes aegypti GN=AAEL009987 PE=4 SV=1
CCG011001.1	1617	0	13	0.001	0.281049844	8.134682205	Up	0.00048	0.000755	Putative uncharacterized protein GLEAN_15470 OS=Tribolium castaneum GN=GLEAN_15470 PE=4 SV=1
CCG027607.1	1902	0	15	0.001	0.275696189	8.106935515	Up	0.00015	0.000236	Threonine--tRNA ligase OS=Wolbachia pipientis wAlbB GN=thrS PE=3 SV=1
CCG027695.1	506	2	657	0.1698408	45.3904898	8.062062764	Up	9.36E-05	0.000153	Predicted protein OS=Nematostella vectensis GN=v1g229551 PE=3 SV=1
CCG021312.1	1662	2	646	0.0517085	13.58787412	8.037703559	Up	9.36E-05	0.000153	AAEL000889-PA (Fragment) OS=Aedes aegypti GN=AAEL000889 PE=4 SV=2
CCG005939.1	701	2	614	0.1225955	30.6196604	7.964408049	Up	9.36E-05	0.000154	-
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
CCG004419.1	1046	2	590	0.0821601	19.71833967	7.906884348	Up	9.36E-05	0.000153	Similar to Aedes aegypti 34 kDa salivary secreted protein 34k-2 OS=Aedes albopictus PE=2 SV=1
CCG027500.1	1100	1	295	0.0390634	9.375174225	7.906884348	Up	0.00059	0.000917	AAEL002875-PA OS=Aedes aegypti GN=AAEL002875 PE=4 SV=1
CCG004788.1	1896	0	13	0.001	0.239692826	7.905042919	Up	0.00048	0.000756	-
CCG013769.1	852	1	288	0.050434	11.81688229	7.872238205	Up	0.00059	0.000919	Putative salivary mucin OS=Aedes albopictus PE=2 SV=1
CCG018500.1	742	2	573	0.1158214	26.99608166	7.864704533	Up	9.36E-05	0.000154	AAEL005916-PA OS=Aedes aegypti GN=AAEL005916 PE=4 SV=1

CCG014537.1	563	2	541	0.1526456	33.59223399	7.781797988	Up	9.36E-05	0.000153	Histone H2B OS=Aedes aegypti GN=AAEL015674 PE=3 SV=1
CCG010648.1	425	1	260	0.1011053	21.38623991	7.724681017	Up	0.00059	0.000919	Histone H2B OS=Aedes aegypti GN=AAEL015674 PE=3 SV=1
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
CCG018777.1	1083	1	250	0.0396766	8.069777648	7.668097488	Up	0.00059	0.000917	Arginine N-methyltransferase 1 OS=Culex quinquefasciatus GN=CpipJ_CPIJ003726 PE=4 SV=1
CCG010863.1	729	2	474	0.1178868	22.73007296	7.591056453	Up	9.36E-05	0.000154	NOS protein OS=Aedes aegypti GN=nos PE=2 SV=1
CCG015658.1	1857	1	227	0.0231393	4.273305777	7.528861691	Up	0.00059	0.000916	AAEL013916-PA OS=Aedes aegypti GN=AAEL013916 PE=4 SV=1
CCG013426.1	729	2	430	0.1178868	20.62010838	7.450506053	Up	9.36E-05	0.000154	AAEL014276-PA OS=Aedes aegypti GN=AAEL014276 PE=4 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
CCG004528.1	828	1	211	0.0518958	8.908449757	7.423412393	Up	0.00059	0.000916	AAEL001690-PA OS=Aedes aegypti GN=AAEL001690 PE=3 SV=1
CCG006667.1	2652	0	13	0.001	0.171364102	7.420921108	Up	0.00048	0.000755	Uncharacterized protein OS=Anopheles darlingi GN=AND_19618 PE=4 SV=1
CCG024178.1	1245	2	403	0.0690277	11.31581168	7.356949232	Up	9.36E-05	0.000153	AAEL015402-PA OS=Aedes aegypti GN=AAEL015402 PE=4 SV=1
CCG007445.1	2793	0	13	0.001	0.162713068	7.34618631	Up	0.00048	0.000755	AAEL006287-PA OS=Aedes aegypti GN=PIWI7 PE=4 SV=1
CCG010647.1	389	2	396	0.2209241	35.58734602	7.331669824	Up	9.36E-05	0.000154	Chromosome undetermined SCAF14565, whole genome shotgun sequence OS=Tetraodon nigroviridis GN=GSTENG00016696001 PE=3 SV=1
CCG014759.1	852	1	197	0.050434	8.083075733	7.324365023	Up	0.00059	0.000918	AAEL002647-PA OS=Aedes aegypti GN=AAEL002647 PE=4 SV=1
CCG005147.1	1662	1	186	0.0258542	3.912298123	7.241472015	Up	0.00059	0.000918	AAEL006333-PA OS=Aedes aegypti GN=AAEL006333 PE=3 SV=1
CCG011089.1	1107	1	186	0.0388164	5.873748401	7.241472015	Up	0.00059	0.000916	FXa-directed anticoagulant OS=Aedes albopictus PE=2 SV=1
CCG023564.1	614	2	367	0.1399666	20.89525664	7.221949457	Up	9.36E-05	0.000153	GK23465 OS=Drosophila willistoni GN=GK23465 PE=4 SV=1
CCG018505.1	1143	2	357	0.0751876	10.91872687	7.182093468	Up	9.36E-05	0.000154	AAEL005938-PA OS=Aedes aegypti GN=AAEL005938 PE=4 SV=1
CCG003975.1	396	1	170	0.1085094	15.00734104	7.11170414	Up	0.00059	0.000917	-
CCG017353.1	270	1	168	0.1591472	21.75181666	7.094630627	Up	0.00059	0.000918	AAEL000323-PA OS=Aedes aegypti GN=AAEL000323 PE=4 SV=1

CCG011996.1	1068	1	167	0.0402338	5.466322304	7.086017496	Up	0.00059	0.000918	AAEL008570-PA OS=Aedes aegypti GN=AAEL008570 PE=3 SV=1
CCG010906.1	417	2	313	0.2060899	26.23966578	6.992332051	Up	9.36E-05	0.000154	Histone H2B OS=Aedes aegypti GN=AAEL015674 PE=3 SV=1
CCG019377.1	954	2	310	0.0900833	11.35960776	6.978437609	Up	9.36E-05	0.000154	AAEL006643-PA OS=Aedes aegypti GN=AAEL006643 PE=4 SV=1
CCG002658.1	768	2	275	0.1119004	12.51761213	6.805601012	Up	9.36E-05	0.000154	AAEL009491-PA OS=Aedes aegypti GN=AAEL009491 PE=4 SV=1
CCG019440.1	1344	1	132	0.0319715	3.433402183	6.746707323	Up	0.00059	0.000917	AAEL008449-PA OS=Aedes aegypti GN=AAEL008449 PE=4 SV=1
CCG011546.1	324	1	130	0.1326226	14.02646908	6.724681017	Up	0.00059	0.000917	-
CCG024710.1	1515	2	245	0.0567257	5.653318686	6.638951143	Up	9.36E-05	0.000154	AAEL002671-PA OS=Aedes aegypti GN=AAEL002671 PE=4 SV=1
CCG000646.1	870	1	119	0.0493905	4.781649352	6.597130967	Up	0.00059	0.000916	Putative secreted protein OS=Aedes aegypti PE=2 SV=1
CCG002191.1	261	2	237	0.32927	31.74372259	6.591056453	Up	9.36E-05	0.000154	AAEL012504-PA OS=Aedes aegypti GN=AAEL012504 PE=4 SV=1
CCG008526.1	483	1	114	0.0889643	8.251021847	6.535203218	Up	0.00059	0.000916	Uncharacterized protein OS=Anopheles darlingi GN=AND_17523 PE=4 SV=1
CCG006107.1	643	1	114	0.066827	6.197890439	6.535203218	Up	0.00059	0.000917	-
CCG001064.1	822	1	108	0.0522746	4.593058262	6.457200706	Up	0.00059	0.000919	Uncharacterized protein OS=Tetrapisispora phaffii (strain ATCC 24235 / CBS 4417 / NBRC 1672 / NRRL Y-8282 / UCD 70-5) GN=TPHA0B00750 PE=4 SV=1
CCG012108.1	372	2	192	0.2310201	18.04298156	6.287275705	Up	9.36E-05	0.000153	AGAP006823-PA OS=Anopheles gambiae GN=AGAP006823 PE=4 SV=1
CCG000295.1	768	1	96	0.0559502	4.369784597	6.287275705	Up	0.00059	0.000917	AAEL004467-PA OS=Aedes aegypti GN=AAEL004467 PE=2 SV=1
CCG002691.1	593	2	190	0.1449232	11.20079694	6.272168812	Up	9.36E-05	0.000154	-
CCG000500.1	723	2	180	0.1188651	8.70330542	6.1941663	Up	9.36E-05	0.000154	AAEL003987-PA OS=Aedes aegypti GN=AAEL003987 PE=4 SV=1
CCG026159.1	753	2	175	0.1141294	8.124433513	6.153524316	Up	9.36E-05	0.000153	AAEL015464-PA OS=Aedes aegypti GN=AAEL015464 PE=3 SV=1
CCG010749.1	1725	2	175	0.04982	3.546491846	6.153524316	Up	9.36E-05	0.000154	Putative salivary secreted protein 62k-3 OS=Aedes albopictus PE=2 SV=1
CCG008249.1	426	2	173	0.2017358	14.19667108	6.136941431	Up	9.36E-05	0.000153	-
CCG024075.1	279	2	170	0.3080268	21.30074212	6.11170414	Up	9.36E-05	0.000153	-

CCG026009.1	294	1	83	0.1461556	9.869173374	6.077352635	Up	0.00059	0.000918	Putative 8 kDa secreted peptide 8kD-2 OS=Aedes albopictus PE=4 SV=1
CCG007962.1	2772	2	162	0.0310027	2.043016175	6.042163207	Up	9.36E-05	0.000154	AAEL009308-PA OS=Aedes aegypti GN=AAEL009308 PE=4 SV=1
CCG004424.1	948	2	162	0.0906534	5.97388274	6.042163207	Up	9.36E-05	0.000154	Putative salivary protein 34k-1 OS=Aedes albopictus PE=2 SV=1
CCG019835.1	810	1	81	0.0530491	3.495827677	6.042163207	Up	0.00059	0.000918	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014520 PE=3 SV=1
CCG018430.1	402	1	80	0.1068899	6.956870999	6.024241299	Up	0.00059	0.000917	AAEL006551-PA OS=Aedes aegypti GN=AAEL006551 PE=4 SV=1
CCG012830.1	1298	1	79	0.0331046	2.127660913	6.006093952	Up	0.00059	0.000916	AAEL003526-PA OS=Aedes aegypti GN=AAEL003526 PE=4 SV=1
CCG025978.1	264	2	156	0.3255283	20.65716355	5.987715423	Up	9.36E-05	0.000154	-
CCG004373.1	885	2	153	0.0971067	6.043634289	5.959701047	Up	9.36E-05	0.000154	-
CCG014082.1	1114	2	149	0.0771449	4.67574797	5.921481724	Up	9.36E-05	0.000154	AAEL004775-PA OS=Aedes aegypti GN=AAEL004775 PE=4 SV=1
CCG015710.1	297	2	147	0.2893585	17.30258143	5.901985549	Up	9.36E-05	0.000154	AAEL007013-PA OS=Aedes aegypti GN=AAEL007013 PE=4 SV=1
CCG006920.1	474	2	146	0.1813069	10.76773926	5.892137763	Up	9.36E-05	0.000154	AAEL004436-PA OS=Aedes aegypti GN=AAEL004436 PE=4 SV=1
CCG028497.1	918	1	69	0.046808	2.627582895	5.810837661	Up	0.00059	0.000917	AAEL012339-PA OS=Aedes aegypti GN=AAEL012339 PE=4 SV=1
CCG008860.1	882	2	137	0.097437	5.430027118	5.800345287	Up	9.36E-05	0.000154	Pyrroline-5-carboxylate reductase OS=Aedes aegypti GN=P5CR-1 PE=2 SV=1
CCG008244.1	1260	2	134	0.0682059	3.71778499	5.768402394	Up	9.36E-05	0.000154	Putative salivary serpin OS=Aedes albopictus PE=2 SV=1
CCG006003.1	945	1	67	0.0454706	2.478523327	5.768402394	Up	0.00059	0.000918	-
CCG017929.1	693	1	67	0.0620054	3.379804536	5.768402394	Up	0.00059	0.000918	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008409 PE=4 SV=1
CCG008173.1	3822	2	133	0.0224855	1.216496811	5.757595639	Up	9.36E-05	0.000154	AAEL002094-PA OS=Aedes aegypti GN=AAEL002094 PE=4 SV=1
CCG002319.1	894	2	133	0.0961292	5.200727976	5.757595639	Up	9.36E-05	0.000154	AGAP005746-PA OS=Anopheles gambiae GN=AGAP005746 PE=4 SV=3
CCG020527.1	414	2	128	0.2075833	10.8083561	5.702313204	Up	9.36E-05	0.000153	-
CCG014538.1	846	1	64	0.0507916	2.64459777	5.702313204	Up	0.00059	0.000916	AAEL006359-PA OS=Aedes aegypti GN=AAEL006359 PE=4 SV=1

CCG009804.1	873	1	64	0.0492208	2.562806086	5.702313204	Up	0.00059	0.000917	AAEL005972-PA OS=Aedes aegypti GN=AAEL005972 PE=4 SV=1
CCG021075.1	453	1	61	0.0948559	4.707405923	5.633050541	Up	0.00059	0.000916	-
CCG025780.1	1164	1	58	0.0369156	1.741907262	5.560294199	Up	0.00059	0.000916	AAEL000130-PA OS=Aedes aegypti GN=AAEL000130 PE=4 SV=1
CCG019655.1	1536	2	115	0.0559502	2.617318899	5.547803255	Up	9.36E-05	0.000154	AAEL002743-PA OS=Aedes aegypti GN=AAEL002743 PE=4 SV=1
CCG005446.1	750	1	57	0.057293	2.656829035	5.535203218	Up	0.00059	0.000917	AAEL002960-PA OS=Aedes aegypti GN=AAEL002960 PE=4 SV=1
CCG015978.1	705	2	112	0.1219	5.553655317	5.509668126	Up	9.36E-05	0.000154	AAEL003906-PA OS=Aedes aegypti GN=AAEL003906 PE=4 SV=1
CCG018104.1	468	1	54	0.0918157	4.03364732	5.457200706	Up	0.00059	0.000918	AAEL009493-PA OS=Aedes aegypti GN=AAEL009493 PE=4 SV=1
CCG005014.1	790	1	53	0.0543921	2.345302113	5.430233658	Up	0.00059	0.000918	-
CCG018637.1	675	2	104	0.1273177	5.386164125	5.402752922	Up	9.36E-05	0.000154	-
CCG015182.1	1257	2	100	0.0683687	2.781088049	5.346169394	Up	9.36E-05	0.000154	AAEL000429-PA OS=Aedes aegypti GN=AAEL000429 PE=4 SV=1
CCG020531.1	582	1	50	0.0738312	3.003288382	5.346169394	Up	0.00059	0.000916	AGAP004106-PA OS=Anopheles gambiae GN=AgaP_AGAP004106 PE=4 SV=5
CCG018339.1	945	1	50	0.0454706	1.849644274	5.346169394	Up	0.00059	0.000915	Putative uncharacterized protein OS=Tribolium castaneum GN=TcasGA2_TC004194 PE=4 SV=1
CCG027277.1	576	1	50	0.0746002	3.034572636	5.346169394	Up	0.00059	0.000918	AAEL004355-PA OS=Aedes aegypti GN=AAEL004355 PE=4 SV=1
CCG025877.1	486	2	98	0.1768302	7.049199843	5.317023048	Up	9.36E-05	0.000153	AAEL009296-PA OS=Aedes aegypti GN=AAEL009296 PE=4 SV=1
CCG025157.1	633	2	98	0.1357654	5.41218187	5.317023048	Up	9.36E-05	0.000154	AGAP010064-PA OS=Anopheles gambiae GN=AGAP010064 PE=4 SV=1
CCG018724.1	1524	1	48	0.0281954	1.101048087	5.287275705	Up	0.00059	0.000916	AAEL004975-PA OS=Aedes aegypti GN=AAEL004975 PE=4 SV=1
CCG027550.2	888	2	95	0.0967787	3.739905736	5.272168812	Up	9.36E-05	0.000153	AAEL008084-PA OS=Aedes aegypti GN=AAEL008084 PE=4 SV=1
CCG020853.1	972	2	95	0.0884151	3.416704006	5.272168812	Up	9.36E-05	0.000154	AAEL009134-PA (Fragment) OS=Aedes aegypti GN=AAEL009134 PE=4 SV=1
CCG004626.1	1344	1	46	0.0319715	1.19648864	5.22587516	Up	0.00059	0.000917	AAEL008449-PA OS=Aedes aegypti GN=AAEL008449 PE=4 SV=1

CCG019520.1	1239	2	88	0.069362	2.482912313	5.161744822	Up	9.36E-05	0.000153	-
CCG005063.1	1096	1	44	0.039206	1.403434469	5.161744822	Up	0.00059	0.000917	-
CCG002818.1	709	1	43	0.0606061	2.120177576	5.128577959	Up	0.00059	0.000916	-
CCG019318.1	471	1	43	0.0912309	3.19151996	5.128577959	Up	0.00059	0.000917	AAEL006670-PA OS=Aedes aegypti GN=AAEL006670 PE=4 SV=1
CCG011361.1	1464	1	43	0.0293509	1.026779987	5.128577959	Up	0.00059	0.000918	AAEL004206-PA OS=Aedes aegypti GN=AAEL004206 PE=4 SV=1
CCG020265.1	336	1	42	0.1278861	4.369784597	5.094630627	Up	0.00059	0.000918	-
CCG005309.1	342	2	78	0.251285	7.972940316	4.987715423	Up	9.36E-05	0.000154	-
CCG004888.2	1446	1	39	0.0297163	0.942858087	4.987715423	Up	0.00059	0.000915	AAEL012230-PA OS=Aedes aegypti GN=AAEL012230 PE=4 SV=1
CCG011919.1	7555	2	77	0.0113752	0.356292166	4.969099745	Up	9.36E-05	0.000153	-
CCG026407.1	1280	1	38	0.0335701	1.037823842	4.950240717	Up	0.00059	0.000915	-
CCG014487.1	354	2	75	0.2427669	7.40641457	4.931131894	Up	9.36E-05	0.000154	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ005832 PE=4 SV=1
CCG014411.1	810	2	75	0.1060981	3.236877479	4.931131894	Up	9.36E-05	0.000153	Uncharacterized protein OS=Anopheles darlingi GN=AND_21799 PE=4 SV=1
CCG028235.1	567	2	74	0.1515687	4.562455875	4.911766569	Up	9.36E-05	0.000154	AAEL007039-PA OS=Aedes aegypti GN=PGRPS5_b PE=4 SV=1
CCG019605.1	1542	1	37	0.0278662	0.838817277	4.911766569	Up	0.00059	0.000915	AAEL011028-PA OS=Aedes aegypti GN=AAEL011028 PE=4 SV=1
CCG014491.1	975	2	73	0.088143	2.617388928	4.892137763	Up	9.36E-05	0.000154	AAEL006329-PA OS=Aedes aegypti GN=AAEL006329 PE=4 SV=1
CCG014322.1	480	1	36	0.0895203	2.621870758	4.872238205	Up	0.00059	0.000915	AAEL009770-PA OS=Aedes aegypti GN=AAEL009770 PE=2 SV=1
CCG024491.1	725	1	36	0.0592686	1.73585926	4.872238205	Up	0.00059	0.000915	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG009757.1	396	2	71	0.2170189	6.267771846	4.852060323	Up	9.36E-05	0.000154	AAEL006197-PA OS=Aedes aegypti GN=AAEL006197 PE=4 SV=1
CCG022958.1	633	1	35	0.0678827	1.932922096	4.831596221	Up	0.00059	0.000915	AAEL000896-PA OS=Aedes aegypti GN=AAEL000896 PE=4 SV=1
CCG015907.1	567	1	35	0.0757844	2.157918319	4.831596221	Up	0.00059	0.000915	AAEL007039-PA OS=Aedes aegypti GN=PGRPS5_b PE=4 SV=1
CCG000691.1	594	1	34	0.0723396	2.000978805	4.789776045	Up	0.00059	0.000915	AAEL003775-PA OS=Aedes aegypti GN=AAEL003775 PE=4 SV=1
CCG019071.1	282	1	34	0.1523749	4.214827696	4.789776045	Up	0.00059	0.000915	AAEL008863-PA OS=Aedes aegypti GN=AAEL008863 PE=4 SV=1

CCG007916.1	2372	1	34	0.0181154	0.501088284	4.789776045	Up	0.00059	0.000915	Putative uncharacterized protein OS=Caenorhabditis remanei GN=CRE_08209 PE=4 SV=1
CCG023021.1	490	1	34	0.0876933	2.425676347	4.789776045	Up	0.00059	0.000915	AAEL010576-PN OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG008028.1	2383	2	65	0.0360636	0.953540911	4.724681017	Up	9.36E-05	0.000154	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG002198.1	561	1	32	0.0765949	1.994055003	4.702313204	Up	0.00059	0.000915	AAEL008337-PA OS=Aedes aegypti GN=AAEL008337 PE=4 SV=1
CCG007624.1	642	2	63	0.1338621	3.430485104	4.679593127	Up	9.36E-05	0.000154	AAEL010375-PA OS=Aedes aegypti GN=AAEL010375 PE=4 SV=1
CCG006728.1	1506	1	31	0.0285324	0.719592683	4.656509514	Up	0.00059	0.000915	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ006292 PE=4 SV=1
CCG022573.1	417	1	31	0.1030449	2.598816738	4.656509514	Up	0.00059	0.000915	AAEL001539-PA (Fragment) OS=Aedes aegypti GN=AAEL001539 PE=4 SV=1
CCG007143.1	1635	2	61	0.0525624	1.304253751	4.633050541	Up	9.36E-05	0.000154	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ018804 PE=4 SV=1
CCG002930.1	204	2	61	0.4212719	10.45321021	4.633050541	Up	9.36E-05	0.000154	AAEL003533-PA OS=Aedes aegypti GN=AAEL003533 PE=4 SV=1
CCG006327.1	537	1	30	0.0800181	1.952976356	4.609203799	Up	0.00059	0.000915	AAEL007972-PA OS=Aedes aegypti GN=AAEL007972 PE=3 SV=1
CCG025091.1	1446	1	30	0.0297163	0.725275452	4.609203799	Up	0.00059	0.000915	AAEL012230-PA OS=Aedes aegypti GN=AAEL012230 PE=4 SV=1
CCG016472.1	873	1	30	0.0492208	1.201315353	4.609203799	Up	0.00059	0.000915	AAEL004584-PA OS=Aedes aegypti GN=AAEL004584 PE=4 SV=1
CCG007299.1	1080	2	59	0.0795736	1.909757713	4.584956253	Up	9.36E-05	0.000154	AAEL004561-PA OS=Aedes aegypti GN=AAEL004561 PE=4 SV=1
CCG022185.1	5617	2	58	0.0152999	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	0.0564648	1.332181375	4.560294199	Up	9.36E-05	0.000153	AAEL015565-PA OS=Aedes aegypti GN=AAEL015565 PE=4 SV=1
CCG000275.1	762	1	29	0.0563907	1.330433106	4.560294199	Up	0.00059	0.000915	-
CCG014797.1	1152	1	29	0.0373001	0.880026065	4.560294199	Up	0.00059	0.000915	AAEL008459-PA OS=Aedes aegypti GN=AAEL008459 PE=4 SV=1
CCG019376.1	612	1	29	0.070212	1.656519651	4.560294199	Up	0.00059	0.000915	AAEL006639-PA OS=Aedes aegypti GN=AAEL006639 PE=4 SV=1

CCG016055.1	968	2	57	0.0887804	2.05849357	4.535203218	Up	9.36E-05	0.000154	AAEL013652-PA OS=Aedes aegypti GN=AAEL013652 PE=3 SV=1
CCG024059.1	2064	2	57	0.0416373	0.965417527	4.535203218	Up	9.36E-05	0.000153	AAEL012229-PA OS=Aedes aegypti GN=AAEL012229 PE=4 SV=1
CCG003644.1	1182	2	57	0.0727068	1.685805225	4.535203218	Up	9.36E-05	0.000153	AAEL014020-PA OS=Aedes aegypti GN=AAEL014020 PE=4 SV=1
CCG021054.1	726	2	56	0.1183739	2.696506197	4.509668126	Up	9.36E-05	0.000154	AAEL012824-PA OS=Aedes aegypti GN=AAEL012824 PE=3 SV=1
CCG002773.1	1101	2	55	0.0780558	1.746326269	4.483672917	Up	9.36E-05	0.000154	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	0.000154	AAEL006747-PA OS=Aedes aegypti GN=AAEL006747 PE=4 SV=1
CCG009644.1	1275	2	54	0.0674035	1.48058584	4.457200706	Up	9.36E-05	0.000154	AAEL014550-PA (Fragment) OS=Aedes aegypti GN=AAEL014550 PE=3 SV=1
CCG024631.1	519	1	27	0.0827933	1.818638676	4.457200706	Up	0.00059	0.000917	AAEL014348-PB OS=Aedes aegypti GN=CASPS8 PE=4 SV=1
CCG003047.1	324	1	27	0.1326226	2.913189731	4.457200706	Up	0.00059	0.000917	-
CCG024719.1	795	2	54	0.1081	2.37452446	4.457200706	Up	9.36E-05	0.000153	AAEL006919-PA OS=Aedes aegypti GN=AAEL006919 PE=3 SV=1
CCG012020.1	381	2	53	0.2255629	4.862962386	4.430233658	Up	9.36E-05	0.000154	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	0.000154	-
CCG022084.1	1464	2	53	0.0587018	1.265566031	4.430233658	Up	9.36E-05	0.000154	AAEL007619-PA OS=Aedes aegypti GN=TOLL5A PE=4 SV=1
CCG013467.1	426	1	26	0.1008679	2.133603747	4.402752922	Up	0.00059	0.000918	-
CCG001719.1	687	2	50	0.1250938	2.544270507	4.346169394	Up	9.36E-05	0.000153	AAEL008482-PA OS=Aedes aegypti GN=AAEL008482 PE=4 SV=1
CCG009344.1	1554	1	25	0.0276511	0.56239184	4.346169394	Up	0.00059	0.000921	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001210 PE=4 SV=1
CCG028585.1	609	1	25	0.0705579	1.435068833	4.346169394	Up	0.00059	0.000922	-
CCG014490.1	497	1	25	0.0864582	1.758464626	4.346169394	Up	0.00059	0.000921	-
CCG015051.1	840	1	25	0.0511544	1.040424904	4.346169394	Up	0.00059	0.000921	AAEL013780-PA OS=Aedes aegypti GN=AAEL013780 PE=4 SV=1
CCG005470.1	319	2	48	0.2694027	5.260179577	4.287275705	Up	9.36E-05	0.000153	-
CCG026325.1	1680	2	48	0.0511544	0.998807908	4.287275705	Up	9.36E-05	0.000153	AAEL014180-PA OS=Aedes aegypti GN=AAEL014180 PE=4 SV=1
CCG022023.1	634	1	24	0.0677756	1.323341707	4.287275705	Up	0.0006	0.000927	-

CCG006414.1	533	1	24	0.0806186	1.574106271	4.287275705	Up	0.0006	0.000927	-
CCG020550.1	513	1	24	0.0837617	1.635474937	4.287275705	Up	0.0006	0.000927	-
CCG005657.1	403	2	48	0.2132493	4.163764975	4.287275705	Up	9.36E-05	0.000153	-
CCG007667.1	3242	1	24	0.0132541	0.258790451	4.287275705	Up	0.0006	0.000927	Elongation factor 1-alpha OS=Aedes aegypti PE=2 SV=1
CCG017399.1	415	2	46	0.2070831	3.874893329	4.22587516	Up	9.36E-05	0.000153	-
CCG022013.1	648	2	46	0.1326226	2.481606067	4.22587516	Up	9.36E-05	0.000153	AAEL011896-PA OS=Aedes aegypti GN=AAEL011896 PE=4 SV=1
CCG006816.1	921	1	23	0.0466555	0.873007998	4.22587516	Up	0.0006	0.000936	AAEL004556-PA OS=Aedes aegypti GN=AAEL004556 PE=4 SV=1
CCG006460.1	528	1	23	0.0813821	1.522803723	4.22587516	Up	0.0006	0.000936	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ003878 PE=4 SV=1
CCG016559.1	2223	2	44	0.0386592	0.691931704	4.161744822	Up	9.36E-05	0.000153	AAEL002752-PA OS=Aedes aegypti GN=AAEL002752 PE=4 SV=1
CCG008314.1	323	2	44	0.2660665	4.762118198	4.161744822	Up	9.36E-05	0.000153	AAEL000278-PA (Fragment) OS=Aedes aegypti GN=AAEL000278 PE=3 SV=1
CCG002259.1	425	1	22	0.1011053	1.809604915	4.161744822	Up	0.00061	0.000953	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	0.00061	0.000953	AAEL006494-PA (Fragment) OS=Aedes aegypti GN=GPRgr45 PE=3 SV=2
CCG000678.1	456	1	22	0.0942319	1.686583528	4.161744822	Up	0.00061	0.000953	AAEL013835-PA OS=Aedes aegypti GN=AAEL013835 PE=4 SV=1
CCG003689.1	933	1	22	0.0460555	0.824310921	4.161744822	Up	0.00061	0.000953	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ008885 PE=4 SV=1
CCG024031.1	633	2	43	0.1357654	2.374732861	4.128577959	Up	9.36E-05	0.000153	AAEL004662-PA OS=Aedes aegypti GN=AAEL004662 PE=4 SV=1
CCG026612.1	2073	2	42	0.0414566	0.708271888	4.094630627	Up	9.36E-05	0.000153	Polyribonucleotide nucleotidyltransferase OS=Wolbachia pipientis wAlbB GN=pnp PE=3 SV=1
CCG011750.1	891	1	21	0.0482264	0.823932449	4.094630627	Up	0.00063	0.000982	-
CCG000734.2	465	1	21	0.092408	1.578760886	4.094630627	Up	0.00063	0.000982	-
CCG005660.1	591	2	41	0.1454137	2.425193482	4.059865208	Up	9.36E-05	0.000153	AAEL013345-PA OS=Aedes aegypti GN=AAEL013345 PE=3 SV=1
CCG009270.1	1695	2	40	0.0507018	0.824974083	4.024241299	Up	9.36E-05	0.000153	AAEL003187-PA (Fragment) OS=Aedes aegypti GN=AAEL003187 PE=4 SV=1
CCG023387.1	780	2	39	0.1101788	1.747913839	3.987715423	Up	9.36E-05	0.000153	-

CCG025537.1	596	2	38	0.1441937	2.228883418	3.950240717	Up	9.36E-05	0.000153	AAEL010576-PI OS=Aedes aegypti GN=AAEL010576 PE=4 SV=1
CCG003231.1	438	2	38	0.1962088	3.032909857	3.950240717	Up	9.36E-05	0.000153	Reverse transcriptase OS=Aedes aegypti GN=pol PE=4 SV=1
CCG015576.1	1591	2	38	0.054016	0.834955699	3.950240717	Up	9.36E-05	0.000153	-
CCG010129.1	609	2	37	0.1411157	2.123901873	3.911766569	Up	9.36E-05	0.000153	AAEL003927-PA OS=Aedes aegypti GN=AAEL003927 PE=4 SV=1
CCG016271.1	1161	2	36	0.0740219	1.083977574	3.872238205	Up	9.36E-05	0.000153	AAEL013852-PA OS=Aedes aegypti GN=AAEL013852 PE=4 SV=1
CCG004412.1	4492	2	36	0.0191317	0.280164284	3.872238205	Up	9.36E-05	0.000153	BEL12_AG transposon polyprotein OS=Anopheles gambiae GN=BEL12-AG transposon PE=4 SV=1
CCG019174.1	399	2	36	0.2153871	3.154130235	3.872238205	Up	9.36E-05	0.000153	AAEL002617-PA OS=Aedes aegypti GN=AAEL002617 PE=4 SV=1
CCG010622.1	1752	2	36	0.0490522	0.718320756	3.872238205	Up	9.36E-05	0.000153	AAEL001986-PA OS=Aedes aegypti GN=AAEL001986 PE=3 SV=1
CCG026332.1	1124	2	35	0.0764586	1.08855844	3.831596221	Up	9.37E-05	0.000153	AAEL007996-PA OS=Aedes aegypti GN=AAEL007996 PE=4 SV=1
CCG004719.1	4358	2	34	0.0197199	0.272735523	3.789776045	Up	9.38E-05	0.000153	Putative uncharacterized protein 882 (3' region) (Fragment) OS=Aedes aegypti PE=4 SV=1
CCG011553.1	489	2	34	0.1757453	2.430636831	3.789776045	Up	9.38E-05	0.000153	AAEL000913-PA OS=Aedes aegypti GN=AAEL000913 PE=4 SV=1
CCG007384.1	1573	2	34	0.0546341	0.755614374	3.789776045	Up	9.38E-05	0.000153	AAEL006747-PA OS=Aedes aegypti GN=AAEL006747 PE=4 SV=1
CCG016872.1	2110	2	34	0.0407296	0.563308725	3.789776045	Up	9.38E-05	0.000153	Uncharacterized protein OS=Anopheles darlingi GN=AND_11211 PE=4 SV=1
CCG020376.1	456	2	34	0.1884637	2.60653818	3.789776045	Up	9.38E-05	0.000153	-
CCG028479.1	1011	2	34	0.0850044	1.175649268	3.789776045	Up	9.38E-05	0.000153	AAEL014410-PA OS=Aedes aegypti GN=AAEL014410 PE=4 SV=1
CCG012240.1	1478	2	33	0.0581458	0.78052986	3.746707323	Up	9.40E-05	0.000154	-
CCG016501.1	1509	2	33	0.0569513	0.764495118	3.746707323	Up	9.40E-05	0.000154	-
CCG013649.1	2117	2	32	0.0405949	0.528419866	3.702313204	Up	9.43E-05	0.000154	AAEL010730-PA OS=Aedes aegypti GN=AAEL010730 PE=4 SV=1
CCG026885.1	1275	2	32	0.0674035	0.877384201	3.702313204	Up	9.43E-05	0.000154	AAEL014814-PA OS=Aedes aegypti GN=AAEL014814 PE=4 SV=1
CCG007709.1	683	2	32	0.1258265	1.637869483	3.702313204	Up	9.43E-05	0.000154	Uncharacterized protein OS=Anopheles darlingi GN=AND_08768 PE=4 SV=1

CCG028224.1	1078	2	32	0.0797212	1.037722502	3.702313204	Up	9.43E-05	0.000154	-
CCG010828.1	447	2	31	0.1922583	2.424399508	3.656509514	Up	9.48E-05	0.000155	AAEL001249-PA OS=Aedes aegypti GN=AAEL001249 PE=4 SV=1
CCG010000.1	552	2	31	0.1556874	1.963236558	3.656509514	Up	9.48E-05	0.000155	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	0.000155	Gambicin OS=Aedes triseriatus PE=4 SV=1
CCG000074.1	1434	2	31	0.0599299	0.755722859	3.656509514	Up	9.48E-05	0.000155	-
CCG012380.1	966	2	30	0.0889643	1.085660769	3.609203799	Up	9.58E-05	0.000156	AAEL014506-PA (Fragment) OS=Aedes aegypti GN=AAEL014506 PE=4 SV=1
CCG020165.1	537	2	30	0.1600363	1.952976356	3.609203799	Up	9.58E-05	0.000156	AAEL006623-PA OS=Aedes aegypti GN=AAEL006623 PE=4 SV=1
CCG020649.1	643	2	29	0.1336539	1.57665634	3.560294199	Up	9.73E-05	0.000159	-
CCG001033.1	642	2	29	0.1338621	1.579112191	3.560294199	Up	9.73E-05	0.000159	-
CCG013766.1	1635	2	29	0.0525624	0.620055062	3.560294199	Up	9.73E-05	0.000159	AAEL011341-PA OS=Aedes aegypti GN=AAEL011341 PE=3 SV=1
CCG014413.1	418	2	29	0.2055968	2.425334991	3.560294199	Up	9.73E-05	0.000159	-
CCG007712.1	1680	2	28	0.0511544	0.582637946	3.509668126	Up	0.0001	0.000163	Chaperone protein DnaK OS=Wolbachia pipientis wAlbB GN=dnaK PE=3 SV=1
CCG017638.1	588	2	28	0.1461556	1.664679846	3.509668126	Up	0.0001	0.000163	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	0.000163	Yellow-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	0.000163	Pol polyprotein (Fragment) OS=Anopheles gambiae PE=4 SV=1
CCG018098.1	1053	2	27	0.0816139	0.896366071	3.457200706	Up	0.0001	0.00017	AAEL011917-PA OS=Aedes aegypti GN=AAEL011917 PE=3 SV=1
CCG011594.1	1560	2	26	0.0550894	0.582637946	3.402752922	Up	0.00011	0.000183	AAEL010194-PA OS=Aedes aegypti GN=AAEL010194 PE=4 SV=1
CCG003349.1	942	2	26	0.0912309	0.964878127	3.402752922	Up	0.00011	0.000183	AAEL005554-PA OS=Aedes aegypti GN=AAEL005554 PE=4 SV=1
CCG012401.1	1655	2	26	0.0519272	0.549193472	3.402752922	Up	0.00011	0.000183	Putative uncharacterized protein (Fragment) OS=Camponotus floridanus GN=EAG_00896 PE=4 SV=1
CCG011245.1	657	2	26	0.1308059	1.383432566	3.402752922	Up	0.00011	0.000183	AAEL010636-PA OS=Aedes aegypti GN=AAEL010636 PE=4 SV=1
CCG018656.1	836	2	25	0.1027984	1.045403014	3.346169394	Up	0.00013	0.000203	AAEL005607-PA OS=Aedes aegypti GN=AAEL005607 PE=3 SV=1

CCG004492.1	838	2	25	0.1025531	1.042908018	3.346169394	Up	0.00013	0.000203	-
CCG018437.1	1701	2	25	0.0505229	0.513790076	3.346169394	Up	0.00013	0.000203	AAEL008252-PA OS=Aedes aegypti GN=AAEL008252 PE=3 SV=1
CCG012172.1	410	2	24	0.2096085	2.046338153	3.287275705	Up	0.00015	0.000238	Predicted protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001647 PE=4 SV=1
CCG015815.1	825	2	24	0.1041691	1.016968052	3.287275705	Up	0.00015	0.000238	AAEL012564-PA OS=Aedes aegypti GN=AAEL012564 PE=3 SV=1
CCG002416.1	1687	2	24	0.0509422	0.497331738	3.287275705	Up	0.00015	0.000238	Putative uncharacterized protein OS=Aedes aegypti PE=4 SV=1
CCG021794.1	297	2	24	0.2893585	2.824911254	3.287275705	Up	0.00015	0.000238	AAEL005936-PA OS=Aedes aegypti GN=AAEL005936 PE=4 SV=1
CCG019917.1	687	2	24	0.1250938	1.221249844	3.287275705	Up	0.00015	0.000238	-
CCG025567.1	648	2	23	0.1326226	1.240803034	3.22587516	Up	0.00018	0.000296	-
CCG016096.1	204	2	23	0.4212719	3.941374342	3.22587516	Up	0.00018	0.000296	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ014565 PE=4 SV=1
CCG006935.1	1371	2	23	0.0626838	0.586462703	3.22587516	Up	0.00018	0.000296	AAEL010721-PA OS=Aedes aegypti GN=AAEL010721 PE=4 SV=1
CCG019990.1	849	2	23	0.1012243	0.947044011	3.22587516	Up	0.00018	0.000296	AAEL013652-PA OS=Aedes aegypti GN=AAEL013652 PE=3 SV=1
CCG028594.1	915	2	23	0.0939229	0.87873264	3.22587516	Up	0.00018	0.000296	AAEL006410-PA OS=Aedes aegypti GN=AAEL006410 PE=4 SV=1
CCG017166.1	1862	2	22	0.0461544	0.413040864	3.161744822	Up	0.00024	0.000392	RNase H and integrase-like protein (Fragment) OS=Bombyx mori GN=P-1 PE=4 SV=1
CCG022134.1	438	2	22	0.1962088	1.75589518	3.161744822	Up	0.00024	0.000392	AGAP000591-PA OS=Anopheles gambiae GN=AgaP_AGAP000591 PE=3 SV=4
CCG016336.1	777	2	21	0.1106042	0.944818291	3.094630627	Up	0.00035	0.000551	AAEL010959-PA OS=Aedes aegypti GN=AAEL010959 PE=3 SV=1
CCG006544.1	363	2	21	0.2367478	2.022379648	3.094630627	Up	0.00035	0.000551	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ001635 PE=4 SV=1
CCG018932.1	1002	2	21	0.0857679	0.732658495	3.094630627	Up	0.00035	0.00055	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ009045 PE=4 SV=1
CCG013642.1	354	2	21	0.2427669	2.07379608	3.094630627	Up	0.00035	0.00055	AAEL009961-PA OS=Aedes aegypti GN=AAEL009961 PE=4 SV=1
CCG025109.1	432	2	21	0.198934	1.699360676	3.094630627	Up	0.00035	0.000551	-
CCG015293.1	1866	2	20	0.0460555	0.374686782	3.024241299	Up	0.00052	0.000812	AAEL004778-PA OS=Aedes aegypti GN=AAEL004778 PE=3 SV=1

CCG006701.1	1521	2	20	0.056502	0.459674908	3.024241299	Up	0.00052	0.000812	AAEL001237-PA (Fragment) OS=Aedes aegypti GN=AAEL001237 PE=4 SV=1
CCG003481.1	1157	2	20	0.0742778	0.604291733	3.024241299	Up	0.00052	0.000812	-
CCG015706.1	477	2	20	0.1801666	1.46575584	3.024241299	Up	0.00052	0.000812	Putative uncharacterized protein OS=Culex quinquefasciatus GN=CpipJ_CPIJ002557 PE=4 SV=1

