

FlyBase gene	vast-tools	Ev	Ensembl	Transcript	Genomic	Strand	Intron length	Intron order
CaMKI	DmeINT0017	FBtr0089067	chr4:446752-	+			69	1
CG11455	DmeINT0002	FBtr0078117	chr2L:155430-	+			137	1
Su(var)205	DmeINT0027	FBtr0079635	chr2L:821080-	-			117	1
CG4709	DmeINT0012	FBtr0079884	chr2L:995450-	+			61	1
CG15362	DmeINT0005	FBtr0077814	chr2L:198150-	+			58	1
mxt	DmeINT0034	FBtr0077409	chr2L:484510-	-			95	3
crol	DmeINT0030	FBtr0080261	chr2L:118030-	-			110	3
hang	DmeINT0032	FBtr0074276	chrX:1643650-	+			83	7
Pmp70	DmeINT0025	FBtr0074795	chrX:1975110-	+			210	1
Ranbp16	DmeINT0025	FBtr0074162	chrX:1592890-	-			410	15
Seipin	DmeINT0027	FBtr0070442	chrX:2620590-	+			231	1
AMPKalpha	DmeINT0000	FBtr0070216	chrX:1375750-	+			106	1
Su(var)2-10	DmeINT0027	FBtr0088578	chr2R:911900-	+			705	7
Su(var)2-10	DmeINT0027	FBtr0088578	chr2R:911980-	+			79	8
tra2	DmeINT0037	FBtr0089617	chr2R:146040-	-			73	1
CG7745	DmeINT0014	FBtr0088119	chr2R:113080-	+			58	1
CG1620	DmeINT0005	FBtr0088983	chr2R:749470-	-			68	2
CG3831	DmeINT0010	FBtr0071993	chr2R:229400-	-			85	1
CG30108	DmeINT0007	FBtr0273298	chr2R:176690-	+			54	1
fl(2)d	DmeINT0031	FBtr0087647	chr2R:138210-	-			2234	1
babo	DmeINT0029	FBtr0088659	chr2R:895580-	+			3248	4
TBPH	DmeINT0028	FBtr0089626	chr2R:238610-	-			803	1
qkr58E-2	DmeINT0035	FBtr0071828	chr2R:224170-	+			56	4
CG6701	DmeINT0014	FBtr0303989	chr2R:139770-	-			64	3
CG18659	DmeINT0006	FBtr0088626	chr2R:912830-	-			60	3
Rpn13	DmeINT0026	FBtr0087610	chr2R:139590-	-			105	3
Git	DmeINT0020	FBtr0088258	chr2R:106450-	-			183	1
CG33145	DmeINT0009	FBtr0088047	chr2R:119810-	-			67	2
prel	DmeINT0035	FBtr0300293	chr2R:942420-	+			65	1
CG7458	DmeINT0014	FBtr0078475	chr3L:219550-	+			520	2
mRpS6	DmeINT0033	FBtr0073288	chr3L:436470-	+			66	2
RhoBTB	DmeINT0026	FBtr0078185	chr3L:203810-	+			254	11
mbf1	DmeINT0033	FBtr0075393	chr3L:165740-	+			100	1
CG7407	DmeINT0014	FBtr0078440	chr3L:218230-	+			62	1
CG11504	DmeINT0002	FBtr0085548	chr3R:299980-	+			64	1
CG1239	DmeINT0003	FBtr0078720	chr3R:573360-	+			70	1
Rad17	DmeINT0025	FBtr0082974	chr3R:146610-	-			58	3
CG7168	DmeINT0014	FBtr0083608	chr3R:182250-	-			78	2
CG5902	DmeINT0013	FBtr0084502	chr3R:239860-	+			85	1
RpL32	DmeINT0026	FBtr0085594	chr3R:300450-	-			200	1
CG13850	DmeINT0004	FBtr0084231	chr3R:224100-	-			175	1
CG11897	DmeINT0002	FBtr0085386	chr3R:291580-	+			151	1
CG31301	DmeINT0008	FBtr0083087	chr3R:153470-	+			56	2
GILT2	DmeINT0020	FBtr0084435	chr3R:236790-	+			66	1
CG18428	DmeINT0006	FBtr0084543	chr3R:241090-	+			67	1

CG8866	DmeINT0015 FBtr0082055 chr3R:93321 -	372	6
Prat	DmeINT0025 FBtr0081764 chr3R:79116 -	80	1
CG5746	DmeINT0013 FBtr0084631 chr3R:24347 -	69	3
CG5514	DmeINT0013 FBtr0085252 chr3R:27942 +	104	4
pygo	DmeINT0035 FBtr0085806 chr3R:31579 -	67	1
bocksbeutel	DmeINT003C FBtr0082105 chr3R:95897 -	100	1
CG31211	DmeINT0008 FBtr0082479 chr3R:11951 +	230	4
CG31211	DmeINT0008 FBtr0082479 chr3R:11953 +	57	8

PIR (Salsa re PIR (Salsa re PIR (Ctrl rep1 PIR (Ctrl rep2 PIR (Salsa) PIR (Ctrl) dPIR (Salsa-(  
0.258291142 0.272977062 0.125857144 0.072214859 0.263048022 0.097250062 0.165797959  
0.490128884 0.444304485 0.405457597 0.318557072 0.466146274 0.359611207 0.106535067  
0.333754507 0.388988045 0.071963382 0.059443600 0.362911004 0.065338462 0.297572542  
0.197652219 0.185633565 0.037678798 0.025158828 0.192933183 0.030989488 0.161943694  
0.155914032 0.162907269 0.056280795 0.048895268 0.160872227 0.051976028 0.108896198  
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0.831680884 0.834300589 0.932586622 0.978831834 0.833513300 0.946294752 -0.11278145  
0.228841547 0.334492234 0.040409572 0.171543076 0.284573269 0.087102298 0.197470971  
0.145135897 0.182770437 0.030155329 0.044090765 0.165406829 0.036240580 0.129166248  
0.113472378 0.119819225 0.008452385 0.005825206 0.112766871 0.006925136 0.105841734  
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0.476518959 0.480252629 0.656393362 0.584094891 0.479198203 0.632360104 -0.15316190  
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0.068681259 0.222835247 0.014402605 0.047611728 0.144566098 0.025231084 0.119335014  
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0.717424543 0.623686715 0.570031177 0.422831121 0.668995965 0.514500741 0.154495223  
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0.535104768 0.617241336 0.290033934 0.413469710 0.580031513 0.348217197 0.231814315  
0.298953800 0.351862309 0.180329214 0.250466235 0.320938074 0.212399347 0.108538727

Pdiff	Predicted cor	Class
0.997	5' UTR	Affected
0.982	5' UTR	Affected
	1 5' UTR	Affected
0.998	ORF disrupti	Affected low PIR
0.994	ORF disrupti	Affected
0.95	ORF disrupti	Affected
0.994	5' UTR	Affected
	1 Alternative p	Less Retained
0.974	5' UTR	Affected
0.995	Alternative p	Affected low PIR
0.994	ORF disrupti	Affected low PIR
	1 5' UTR	Affected
0.994	ORF disrupti	Less Retained
0.997	Alternative p	Less Retained
0.999	5' UTR	Affected
0.907	Alternative p	Affected low PIR
0.992	Alternative p	Affected
0.999	ORF disrupti	Affected low PIR
0.997	Alternative p	Affected
0.907	ORF disrupti	Affected
0.997	ORF disrupti	Affected
0.913	5' UTR	Affected
0.996	3' UTR	Affected
0.931	ORF disrupti	Affected
0.991	Alternative p	Affected
	1 Alternative p	Affected
0.95	5' UTR	Affected
0.968	Alternative p	Less Retained
0.947	5' UTR	Affected
0.98	ORF disrupti	Affected
0.996	Alternative p	Affected low PIR
0.9	ORF disrupti	Affected
	1 5' UTR	Affected
0.999	Alternative p	Affected low PIR
	1 5' UTR	Affected
	1 5' UTR	Affected
0.994	Alternative p	Affected low PIR
	1 5' UTR	Affected
0.971	5' UTR	Affected
0.98	5' UTR	Affected
0.992	5' UTR	Affected low PIR
0.99	5' UTR	Affected
	1 ORF disrupti	Affected
	1 Alternative p	Affected low PIR
	1 Alternative p	Affected

0.928	Alternative p	Affected
0.989	5' UTR	Affected
	1 Alternative p	Affected
0.984	3' UTR	Less Retained
0.93	5' UTR	Less Retained
0.964	Alternative p	Affected
0.982	ORF disrupti	Affected
0.918	Alternative p	Affected