

[illegible][illegible][illegible]

	PASS/FAIL
crit.star	0
crit.loop	0
crit.mor	0
crit.half	0
crit.total	1
crit.pairing	0
crit.top3	1
crit.tgtop3	1
crit.uri	0
crit.back	0
rescue.total	1
rescue.dominant	1
rescue.known	0
rescue.confident	3
rescue.candidate	0



Flybase annotation

intergenic

No Repeatable elements found

Sense Strand Reads

hide 3p reads | show mid mismatch reads

Anti-sense strand reads

CTTAGACACGCGAGATAGGGTCAATTGCGTGCAGAAATTCCTTGGCAATAAGGGTAATGCACGACAAGCTCTCAGACACCTGCACAGGTTTAAACAATGTCGGTGACGCTAAGGCCCGTCAACTTTCACATTATGCCCTAGGGGTTTTGTGTAGTAGACGCGAAACAGTGGCGGGTGCTCT															V055	V039	M058	V105
*****(((.....(((.....(((.....(((.....(((.....)))))))).....)))).....))))*****										Read #	Hit	Total			head	embryo	embryo	male
										size	Mismatch	Count	Norm	Total				body

Show Alignment With Reads

Re-alignment of all predicted orthologs

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