

SNiP Acard

Block annotations

Block info

genomic range	chr17:34,475,131-34,475,131 <i>e!</i>
block size	1 bp
variant count	1 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	-1.953	gene(s) hit or close-by	CTB-91J4.1 <i>e!</i>
phastCons	0	eQTL gene(s)	-
GERP++	-2.76	potentially regulated gene(s)	-
CADD score	0.344	disease gene(s)	-

Putative effect on transcript

Intron variant

gene	affected transcript	RefSeq id	protein	variant(s)
CTB-91J4.1 <i>e!</i>	ENST00000617914 <i>e!</i>	?	?	1

