Locus 284 (rs3213636)

SNiPAcard

Block annotations

Block info

genomic range	chr17:34,475,131-34,475,131
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
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	ENST00000617914	?	?	1