

SNiPacard

Block annotations

Block info


genomic range	chr6:31,572,333-31,572,333 <i>e!</i>
block size	1 bp
variant count	1 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	-0.269	gene(s) hit or close-by	-
phastCons	0	eQTL gene(s)	XXbac-BPG181B23.7 <i>e!</i>
GERP++	-1.97	potentially regulated gene(s)	-
CADD score	0.854	disease gene(s)	-

Direct effect on regulation

cis-eQTL

gene	transcript	probe	tissue	min(statistic) (type)	source	variant(s)
XXbac-BPG181B23.7 <i>e!</i>	?	ENSG00000272221 <i>e!</i>	skeletal muscle	3.60×10 ⁻⁶ (p-value)	GTEx Portal V6 	1

