Locus 132 (rs2736186)

SNiPAcard

Block annotations

Block info

genomic range	chr6:31,572,333-31,572,333
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
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	?	ENSG00000272221	skeletal muscle	3.60×10^{-6} (p-value)	GTEx Portal V6	1