Locus 276 (rs259842)

SNiPAcard

Block annotations

Block info

genomic range	chr2:180,738,840-180,739,450
block size	611 bp
variant count	2 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = 0.499 [0.068 – 0.929]	**gene(s) hit or close-by**	–
phastCons	μ = 0.102 [0 – 0.203]	**eQTL gene(s)**	ZNF385B
GERP++	μ = 0.581 [-0.0672 – 1.23]	**potentially regulated gene(s)**	–
CADD score	μ = 5.518 [4.902 – 6.135]	**disease gene(s)**	–

Direct effect on regulation

cis-eQTL

gene	transcript	probe	tissue	min(statistic) (type)	source	variant(s)
ZNF385B	?	ENSG00000144331	tibial artery	2.09×10^{-6} (p-value)	GTEx Portal V6	2