Locus 278 (rs6603134)

SNiPAcard

Block annotations

Block info

genomic range	chr19:8,105,520-8,107,703
block size	2,184 bp
variant count	4 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = -1.185 [-2.036 – 0.169]	**gene(s) hit or close-by**	ELAVL1
phastCons	μ = 0.012 [0.002 – 0.024]	**eQTL gene(s)**	–
GERP++	μ = -0.042 [-0.167 – 0]	**potentially regulated gene(s)**	–
CADD score	μ = 1.701 [0.781 – 3.285]	**disease gene(s)**	–

Putative effect on transcript

Intron variant

gene	affected transcript	RefSeq id	protein	variant(s)
ELAVL1	ENST00000351593	?	ENSP00000264073	4