Locus 148 (rs2361355)

SNiPAcard

Block annotations

Block info

genomic range	chr12:94,621,186-94,630,029
block size	8,844 bp
variant count	11 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = -0.564 [-1.928 – 1.142]	**gene(s) hit or close-by**	PLXNC1
phastCons	μ = 0.166 [0 – 0.99]	**eQTL gene(s)**	–
GERP++	μ = -0.797 [-3.36 – 3.49]	**potentially regulated gene(s)**	–
CADD score	μ = 4.859 [0.817 – 16.82]	**disease gene(s)**	–

Putative effect on regulation

Variation proximal to gene

gene	variant type	min(distance)	transcript	RefSeq id	protein	variant(s)
PLXNC1	downstream gene variant	386	ENST00000546659	?	?	7

Putative effect on transcript

Intron variant

gene	affected transcript	RefSeq id	protein	variant(s)
PLXNC1	ENST00000258526	NM_005761.2	ENSP00000258526	11

Non-coding exon variant

gene	affected transcript	RefSeq id	variant(s)
PLXNC1	ENST00000546659	?	1