Locus 369 (rs16971895)

SNiPAcard

Block annotations

Block info

genomic range	chr17:34,383,107-34,391,237
block size	8,131 bp
variant count	5 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = -0.784 [-2.213 – 0.281]	**gene(s) hit or close-by**	CCL18
phastCons	μ = 0.008 [0 – 0.027]	**eQTL gene(s)**	–
GERP++	μ = -2.376 [-4.62 – 0.235]	**potentially regulated gene(s)**	–
CADD score	μ = 1.184 [0.023 – 3.986]	**disease gene(s)**	–

Putative effect on regulation

Variation proximal to gene

gene	variant type	min(distance)	transcript	RefSeq id	protein	variant(s)
CCL18	upstream gene variant	403	ENST00000616054	NM_002988.2	ENSP00000479955	3