Locus 331 (rs2989736)

SNiPAcard

Block annotations

Block info

genomic range	chr9:137,797,175-137,797,175
block size	1 bp
variant count	1 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	-3.233	**gene(s) hit or close-by**	FCN1
phastCons	0	**eQTL gene(s)**	–
GERP++	-1.67	**potentially regulated gene(s)**	–
CADD score	0.312	**disease gene(s)**	–

Putative effect on regulation

Variation proximal to gene

gene	variant type	min(distance)	transcript	RefSeq id	protein	variant(s)
FCN1	downstream gene variant	4256	ENST00000371806	NM_002003.3	ENSP00000360871	1
FCN1	downstream gene variant	561	ENST00000616356	?	ENSP00000479379	1