Locus 362 (rs9534338)

SNiPAcard

Block annotations

Block info

genomic range	chr13:46,706,598-46,723,294
block size	16,697 bp
variant count	3 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = -0.108 [-0.764 – 0.273]	**gene(s) hit or close-by**	LCP1
phastCons	μ = 0.001 [0 – 0.002]	**eQTL gene(s)**	–
GERP++	μ = 0.349 [-1.92 – 3.66]	**potentially regulated gene(s)**	–
CADD score	μ = 5.530 [3.363 – 9.165]	**disease gene(s)**	–

Putative effect on regulation

Variation proximal to gene

gene	variant type	min(distance)	transcript	RefSeq id	protein	variant(s)
LCP1	downstream gene variant	3622	ENST00000416500	?	ENSP00000408052	1

Putative effect on transcript

Intron variant

gene	affected transcript	RefSeq id	protein	variant(s)
LCP1	ENST00000494531	?	?	1
LCP1	ENST00000323076	NM_002298.4	ENSP00000315757	3
LCP1	ENST00000398576	?	ENSP00000381581	3
LCP1	ENST00000469227	?	?	1