Locus 373 (rs854658)

SNiPAcard

Block annotations

Block info

genomic range	chr17:34,352,666-34,352,666
block size	1 bp
variant count	1 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	-0.07	gene(s) hit or close-by	–
phastCons	0.16	eQTL gene(s)	CCL14, CCL15-CCL14, CCL23
GERP++	-2	potentially regulated gene(s)	–
CADD score	1.631	disease gene(s)	–

Direct effect on regulation

cis-eQTL

gene	transcript	probe	tissue	min(statistic) (type)	source	variant(s)
CCL23	ENST00000613876	ILMN_1764030	skin	3.00×10^{-5} (p-value)	MuTHER consortium	1
CCL23	ENST00000615050					
CCL23	ENST00000613876	ILMN_1686109	skin	2.01×10^{-5} (p-value)	MuTHER consortium	1
CCL23	ENST00000615050		adipocyte	2.10×10^{-5} (p-value)	MuTHER consortium	1
CCL23	ENST00000612516					
CCL14	ENST00000622526	ILMN_1740609	adipocyte	6.68×10^{-8} (p-value)	MuTHER consortium	1
CCL15-CCL14	ENST00000616694					
CCL15-CCL14	ENST00000610751					
CCL14	ENST00000618404					
CCL14	ENST00000614009					
CCL14	ENST00000620991					