Locus 418 (rs11072996)

SNiPAcard

Block annotations

Block info

genomic range	chr15:81,529,410-81,529,410
block size	1 bp
variant count	1 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	-0.006	gene(s) hit or close-by	IL16
phastCons	0	eQTL gene(s)	–
GERP++	-1.53	potentially regulated gene(s)	–
CADD score	3.993	disease gene(s)	–

Putative effect on transcript

Intron variant

gene	affected transcript	RefSeq id	protein	variant(s)
IL16	ENST00000560241	?	ENSP00000452738	1
IL16	ENST00000394660	NM_001172128.1	ENSP00000378155	1
IL16	ENST00000302987	NM_172217.3	ENSP00000302935	1
IL16	ENST00000559383	?	ENSP00000453250	1
IL16	ENST00000360547	?	ENSP00000456972	1