Locus 425 (rs41265930)

SNiPAcard

Block annotations

Block info

genomic range	chr6:160,964,135-160,964,135
block size	1 bp
variant count	1 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	1.009	gene(s) hit or close-by	LPA
phastCons	0.012	eQTL gene(s)	–
GERP++	0.476	potentially regulated gene(s)	–
CADD score	2.631	disease gene(s)	–

Putative effect on transcript

Intron variant

gene	affected transcript	RefSeq id	protein	variant(s)
LPA	ENST00000316300	NM_005577.2	ENSP00000321334	1