

Block annotations

Block info

genomic range	chr21:28,346,546-28,346,546 <i>e!</i>
block size	1 bp
variant count	1 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	-1.027	gene(s) hit or close-by	-
phastCons	0	eQTL gene(s)	-
GERP++	-1.07	potentially regulated gene(s)	-
CADD score	1.256	disease gene(s)	-

