SNiPAcard

Block annotations

Block info

genomic range	chr10:48,005,225-48,039,886
block size	34,662 bp
variant count	5 variants

Basic features

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