SNiPAcard

Block annotations

Block info

genomic range	chr10:101,786,600-101,792,462
block size	5,863 bp
variant count	6 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = -0.834 [-2.774 – 0.224]	**gene(s) hit or close-by**	–
phastCons	μ = 0.109 [0.004 – 0.353]	**eQTL gene(s)**	–
GERP++	μ = -0.108 [-0.317 – 0.113]	**potentially regulated gene(s)**	–
CADD score	μ = 2.163 [0.156 – 7.061]	**disease gene(s)**	–