SNiPAcard

Block annotations

Block info

genomic range	chr7:82,976,912-82,978,643
block size	1,732 bp
variant count	3 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = 0.437 [0.109 – 0.961]	**gene(s) hit or close-by**	–
phastCons	μ = 0.009 [0.001 – 0.023]	**eQTL gene(s)**	–
GERP++	μ = 0.500 [-0.0267 – 1.41]	**potentially regulated gene(s)**	–
CADD score	μ = 2.549 [0.653 – 4.433]	**disease gene(s)**	–