Locus 359 (rs35871176)

SNiPAcard

Block annotations

Block info

genomic range	chr17:34,366,820-34,375,832
block size	9,013 bp
variant count	4 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = 0.167 [-0.156 – 0.434]	**gene(s) hit or close-by**	–
phastCons	μ = 0.208 [0.013 – 0.709]	**eQTL gene(s)**	–
GERP++	μ = -0.199 [-0.794 – 0]	**potentially regulated gene(s)**	–
CADD score	μ = 2.542 [0.796 – 5.671]	**disease gene(s)**	–