Locus 381 (rs17660992)

SNiPAcard

Block annotations

Block info

genomic range	chr19:52,163,948-52,165,394
block size	1,447 bp
variant count	2 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = -0.183 [-0.401 – 0.036]	**gene(s) hit or close-by**	–
phastCons	μ = 0.000 [0 – 0]	**eQTL gene(s)**	–
GERP++	μ = -0.621 [-1.34 – 0.0977]	**potentially regulated gene(s)**	–
CADD score	μ = 6.307 [4.247 – 8.367]	**disease gene(s)**	–