Locus 361 (rs12411860)

SNiPAcard

Block annotations

Block info

genomic range	chr10:52,032,893-52,043,336
block size	10,444 bp
variant count	19 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = -0.432 [-3.86 – 0.643]	**gene(s) hit or close-by**	–
phastCons	μ = 0.010 [0 – 0.177]	**eQTL gene(s)**	–
GERP++	μ = -0.877 [-5.99 – 3.01]	**potentially regulated gene(s)**	–
CADD score	μ = 2.725 [0.089 – 7.564]	**disease gene(s)**	–