

Block annotations

Block info

genomic range	chr19:52,163,948-52,165,394 <i>e!</i>
block size	1,447 bp
variant count	2 variants

Basic features

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

