SNiPAcard

Block annotations

Block info

genomic range	chr15:60,476,248-60,519,685
block size	43,438 bp
variant count	14 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = -0.168 [-1.523 – 1.224]	**gene(s) hit or close-by**	–
phastCons	μ = 0.005 [0 – 0.032]	**eQTL gene(s)**	–
GERP++	μ = -0.311 [-5.42 – 3.25]	**potentially regulated gene(s)**	–
CADD score	μ = 2.754 [0.377 – 7.678]	**disease gene(s)**	–