SNiPAcard

Block annotations

Block info

genomic range	chr9:120,327,400-120,341,514
block size	14,115 bp
variant count	3 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	μ = 0.006 [-0.331 – 0.323]	**gene(s) hit or close-by**	–
phastCons	μ = 0.332 [0 – 0.99]	**eQTL gene(s)**	–
GERP++	μ = 0.340 [-1.25 – 2.12]	**potentially regulated gene(s)**	ASTN2
CADD score	μ = 5.697 [0.778 – 14.58]	**disease gene(s)**	–

Putative effect on regulation

ENCODE promoter-associated distal DHS (Enhancer)

SNiPA enhancer id	variant(s)	associated SNiPA promoter id	associated gene(s)
ENCE00000552648	1	ENCP00000059825	ASTN2