SNiPAcard

Block annotations

Block info

genomic range	chr11:120,159,052-120,159,052
block size	1 bp
variant count	1 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	1.227	**gene(s) hit or close-by**	POU2F3
phastCons	0.117	**eQTL gene(s)**	–
GERP++	0.843	**potentially regulated gene(s)**	–
CADD score	2.745	**disease gene(s)**	–

Putative effect on transcript

Intron variant

gene	affected transcript	RefSeq id	protein	variant(s)
POU2F3	ENST00000260264	NM_001244682.1	ENSP00000260264	1
POU2F3	ENST00000533620	?	ENSP00000435738	1
POU2F3	ENST00000543440	NM_014352.3	ENSP00000441687	1