SNiPAcard

Block annotations

Block info

genomic range	chr6:160,519,276-160,519,276
block size	1 bp
variant count	1 variants

Basic features

Conservation/deleteriousness		Linked genes	
phyloP	-0.576	**gene(s) hit or close-by**	IGF2R, RP11-288H12.3
phastCons	0	**eQTL gene(s)**	–
GERP++	-2.07	**potentially regulated gene(s)**	–
CADD score	0.968	**disease gene(s)**	–

Trait annotations

Variant association

trait	min(p-value)	source DB	source entry/link	variant(s)
Lipoproteins	4.40×10-5	dbGaP	pha003079 dbGaP	1

Putative effect on regulation

Variation proximal to gene

gene	variant type	min(distance)	transcript	RefSeq id	protein	variant(s)
RP11-288H12.3	downstream gene variant	2032	ENST00000569097	?	?	1
RP11-288H12.3	downstream gene variant	4251	ENST00000366273	?	?	1

Putative effect on transcript

Intron variant

gene	affected transcript	RefSeq id	protein	variant(s)
IGF2R	ENST00000356956	NM_000876.2	ENSP00000349437	1
IGF2R	ENST00000475584	?	?	1