

# SNiPacard

## Block annotations

### Block info

genomic range	chr15:81,529,410-81,529,410 <i>e!</i>
block size	1 bp
variant count	1 variants

### Basic features

Conservation/deleteriousness		Linked genes	
phyloP	-0.006	gene(s) hit or close-by	IL16 <i>e!</i>
phastCons	0	eQTL gene(s)	-
GERP++	-1.53	potentially regulated gene(s)	-
CADD score	3.993	disease gene(s)	-

## Putative effect on transcript

### Intron variant

gene	affected transcript	RefSeq id	protein	variant(s)
IL16 <i>e!</i>	ENST00000560241 <i>e!</i>	?	ENSP00000452738 <i>e!</i>	1
IL16 <i>e!</i>	ENST00000394660 <i>e!</i>	NM_001172128.1	ENSP00000378155 <i>e!</i>	1
IL16 <i>e!</i>	ENST00000302987 <i>e!</i>	NM_172217.3	ENSP00000302935 <i>e!</i>	1
IL16 <i>e!</i>	ENST00000559383 <i>e!</i>	?	ENSP00000453250 <i>e!</i>	1
IL16 <i>e!</i>	ENST00000360547 <i>e!</i>	?	ENSP00000456972 <i>e!</i>	1

