

# SNiPACard

## Block annotations

### Block info

genomic range	chr10:43,435,628-43,447,847 <i>e!</i>
block size	12,220 bp
variant count	12 variants

### Basic features

Conservation/deleteriousness		Linked genes	
phyloP	$\mu = -0.714$ [-2.896 – 0.183]	gene(s) hit or close-by	-
phastCons	$\mu = 0.058$ [0 – 0.377]	eQTL gene(s)	RET <i>e!</i>
GERP++	$\mu = -2.080$ [-8.59 – 0.481]	potentially regulated gene(s)	AL022344.5 <i>e!</i> , CCNYL2 <i>e!</i> , RET <i>e!</i> , RET <i>e!</i> , RP11-124011.1 <i>e!</i>
CADD score	$\mu = 3.098$ [0.535 – 14.03]	disease gene(s)	RET <i>e!</i>

## Trait annotations

### Disease gene annotation

gene	trait	source DB	source entry/link
RET <i>e!</i>	PHEOCHROMOCYTOMAPHEOCHROMOCYTOMA, SUSCEPTIBILITY TO	OMIM	MIM:171300 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	CENTRAL HYPOVENTILATION SYNDROME, CONGENITAL	OMIM	MIM:209880 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	MULTIPLE ENDOCRINE NEOPLASIA, TYPE IIA	OMIM	MIM:171400 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	THYROID CARCINOMA, FAMILIAL MEDULLARY	OMIM	MIM:155240 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	MULTIPLE ENDOCRINE NEOPLASIA, TYPE IIB	OMIM	MIM:162300 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	COLORECTAL CANCER	OMIM	MIM:114500 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	HIRSCHSPRUNG DISEASE, SUSCEPTIBILITY TO, 1	OMIM	MIM:142623 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	THYROID CARCINOMA, PAPILLARY	OMIM	MIM:188550 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	RENAL AGENESIS	DECIPHER	MIM:191830 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	Multiple endocrine neoplasia IIB	DECIPHER	MIM:162300 <i>OMIM</i> <sup>®</sup>
RET <i>e!</i>	Unilateral renal dysplasia	OrphaNet	OrphaNet:93172 <i>orphanet</i>
RET <i>e!</i>	Bilateral renal dysplasia	OrphaNet	OrphaNet:93173 <i>orphanet</i>
RET <i>e!</i>	Haddad syndrome	OrphaNet	OrphaNet:99803 <i>orphanet</i>
RET <i>e!</i>	Hirschsprung disease	OrphaNet	OrphaNet:388 <i>orphanet</i>
RET <i>e!</i>	Multiple endocrine neoplasia type 2A	OrphaNet	OrphaNet:247698 <i>orphanet</i>
RET <i>e!</i>	Papillary or follicular thyroid carcinoma	OrphaNet	OrphaNet:146 <i>orphanet</i>
RET <i>e!</i>	Multiple endocrine neoplasia type 2B	OrphaNet	OrphaNet:247709 <i>orphanet</i>
RET <i>e!</i>	Familial medullary thyroid carcinoma	OrphaNet	OrphaNet:99361 <i>orphanet</i>
RET <i>e!</i>	Bilateral renal agenesis	OrphaNet	OrphaNet:1848 <i>orphanet</i>

## Direct effect on regulation

### cis-eQTL

gene	transcript	probe	tissue	min(statistic) (type)	source	variant(s)
RET <i>e!</i>	?	ENSG00000165731 <i>e!</i>	unexposed skin	1.48×10 <sup>-7</sup> (p-value)	GTEx Portal V6 <i>mq</i>	12

## Putative effect on regulation

### ENCODE promoter-associated distal DHS (Enhancer)

SNiPA enhancer id	variant(s)	associated SNiPA promoter id	associated gene(s)
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ENCE00000072563 <i>e!</i>	1	ENCP00000008221	AL022344.5 <i>e!</i>
		ENCP00000008225	RP11-124011.1 <i>e!</i>
		ENCP00000008213	CCNYL2 <i>e!</i>
		ENCP00000008212	CCNYL2 <i>e!</i>
		ENCP00000008227	RET <i>e!</i>
			RET <i>e!</i>

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