

Rcn1 Cas9-KO Strategy

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Project Overview

Project Name

Rcn1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rcn1* gene. The schematic diagram is as follows:



- The *Rcn1* gene has 2 transcripts. According to the structure of *Rcn1* gene, exon2-exon5 of *Rcn1-201* (ENSMUST00000006128.6) transcript is recommended as the knockout region. The region contains 634bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rcn1* gene. The brief process is as follows: CRISPR/Cas9 system v

- The *Rcn1* gene is located on the Chr2. If the knockout mice are crossed with other mice strains to obtain double gene positive homozygous mouse offspring, please avoid the two genes on the same chromosome.
- This Strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risk of the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

Rcn1 reticulocalbin 1 [Mus musculus (house mouse)]

Gene ID: 19672, updated on 31-Jan-2019

Summary



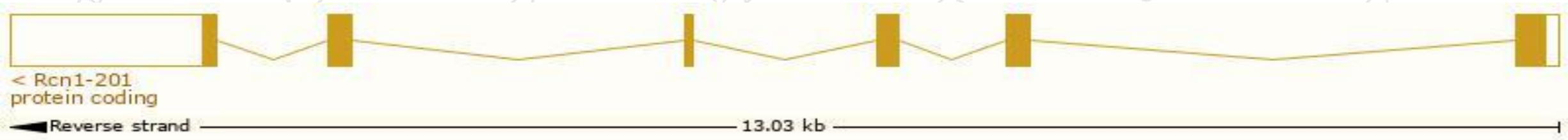
Official Symbol	Rcn1 provided by MGI
Official Full Name	reticulocalbin 1 provided by MGI
Primary source	MGI:MGI:104559
See related	Ensembl:ENSMUSG000000005973
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Rcn
Expression	Broad expression in limb E14.5 (RPKM 50.5), adrenal adult (RPKM 36.1) and 22 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

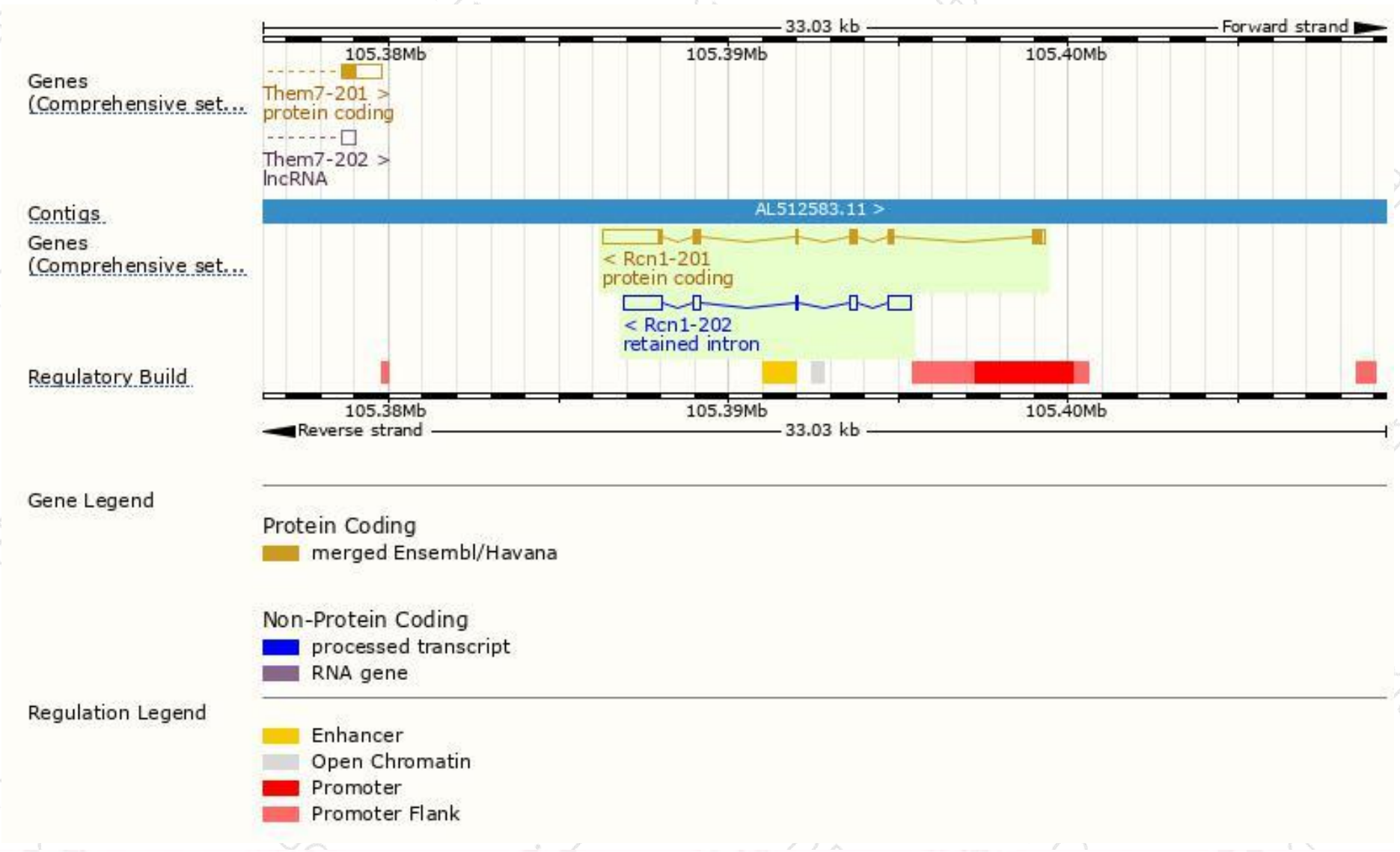
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rcn1-201	ENSMUST00000006128.6	2720	325aa	Protein coding	CCDS16498	Q05186	TSL:1 GENCODE basic APPRIS P1
Rcn1-202	ENSMUST00000127019.1	2208	No protein	Retained intron	-	-	TSL:2

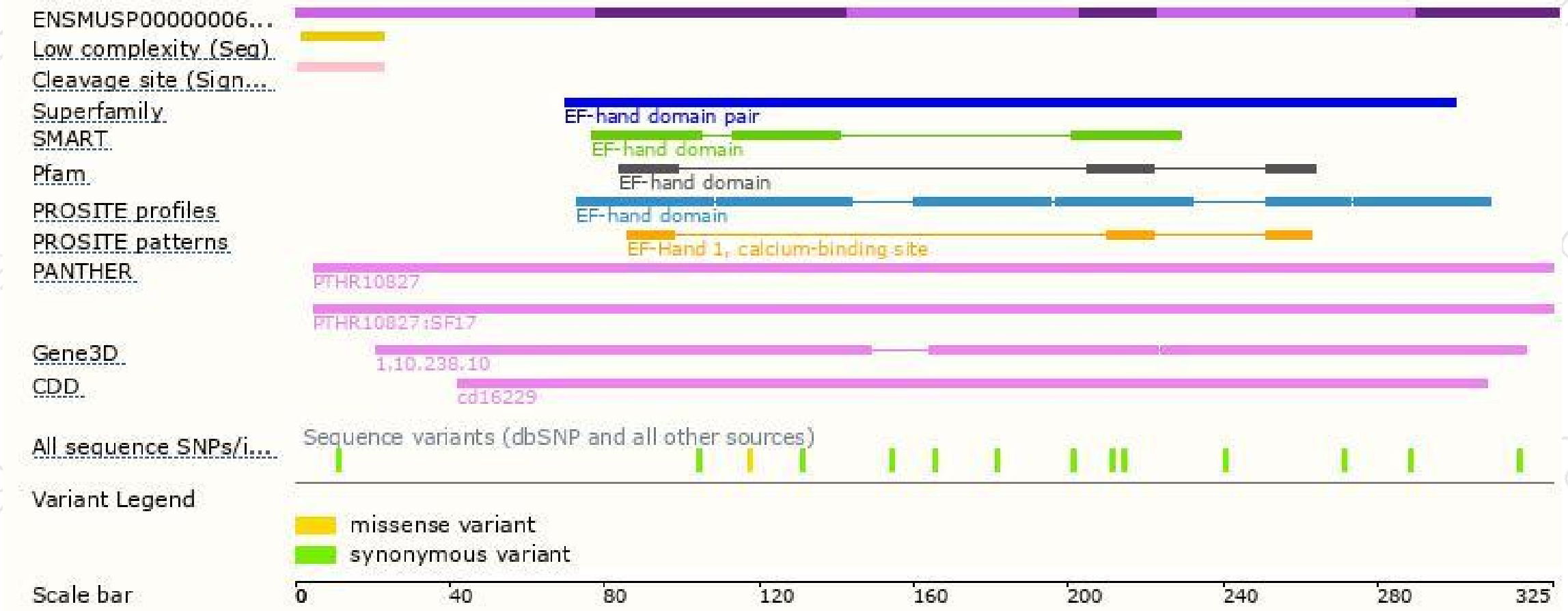
The strategy is based on the design of *Rcn1-201* transcript,The transcription is shown below



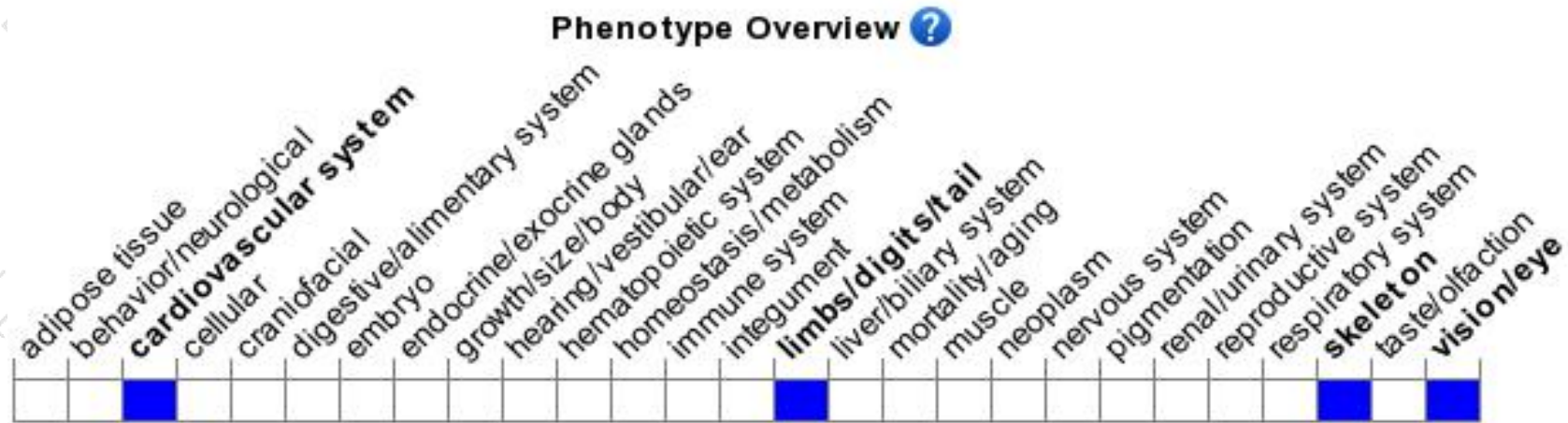
Genomic location distribution



Protein domain



Mouse phenotype description(MGI)



Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).

If you have any questions, you are welcome to inquire.

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