

Rnf169 Cas9-KO Strategy

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Project Overview

Project Name

Rnf169

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rnfl69* gene. The schematic diagram is as follows:



- The *Rnfl69* gene has 2 transcripts. According to the structure of *Rnfl69* gene, exon2 of *Rnfl69-201* (ENSMUST00000080817.5) transcript is recommended as the knockout region. The region contains 74bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rnfl69* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Rnfl69* gene is located on the Chr7. 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.
- The N-terminal of *Rnfl69* gene will remain 154aa, it may remain the partial function of *Rnfl69* gene.
- 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)

Rnf169 ring finger protein 169 [Mus musculus (house mouse)]

Gene ID: 108937, updated on 31-Jan-2019

Summary



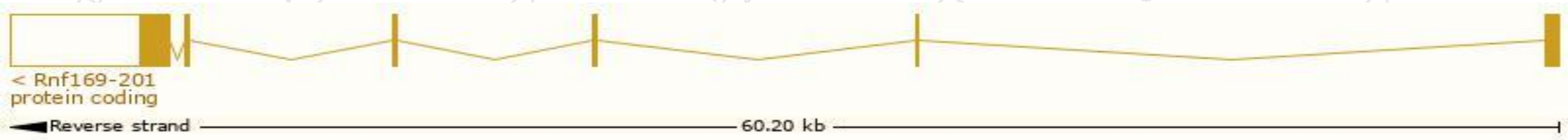
Official Symbol	Rnf169 provided by MGI
Official Full Name	ring finger protein 169 provided by MGI
Primary source	MGI:MGI:1920257
See related	Ensembl:ENSMUSG00000058761
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2900057K09Rik
Expression	Ubiquitous expression in thymus adult (RPKM 1.9), CNS E11.5 (RPKM 1.7) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

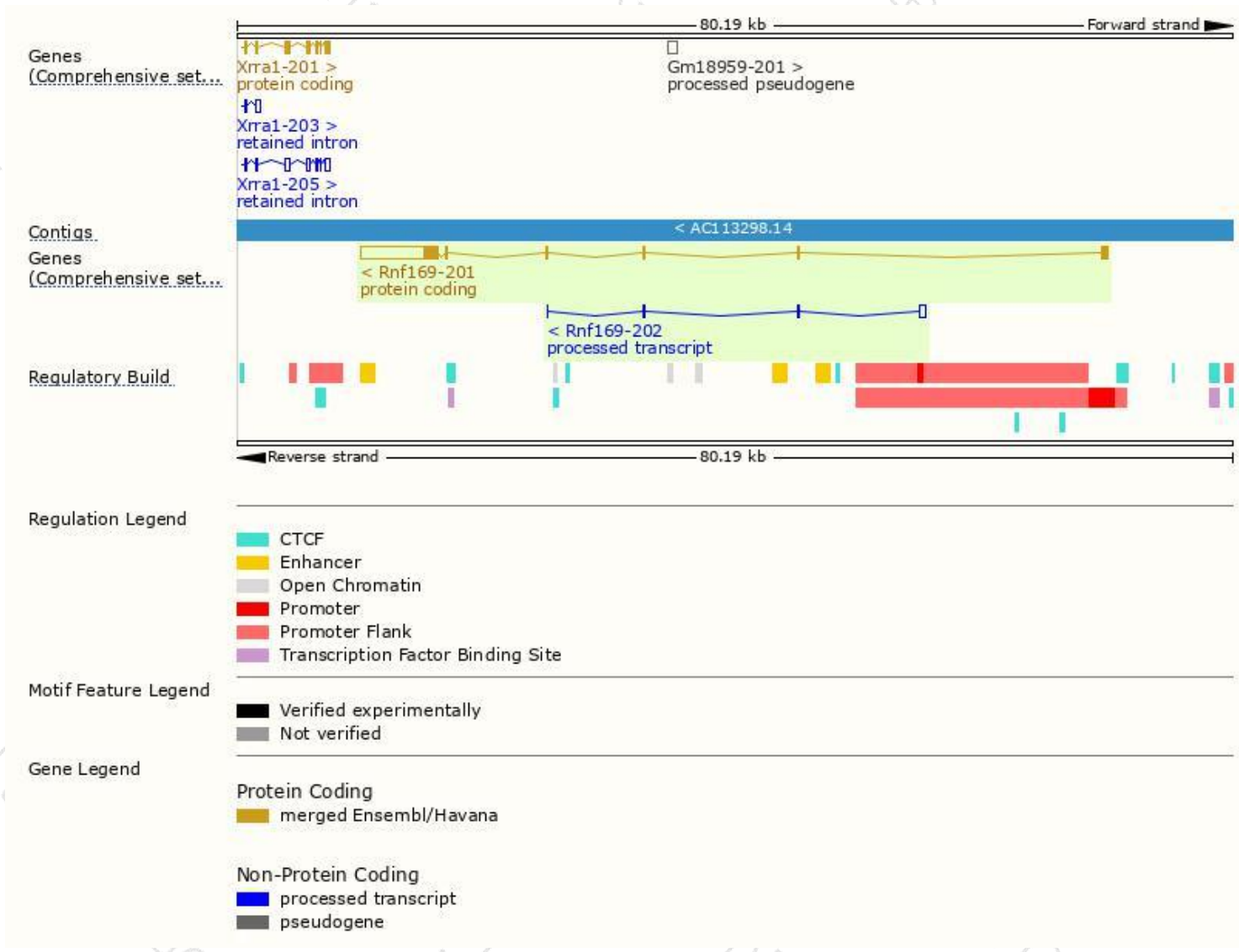
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rnf169-201	ENSMUST00000080817.5	7147	694aa	Protein coding	CCDS52322	E9Q7F2	TSL:5 GENCODE basic APPRIS P1
Rnf169-202	ENSMUST00000174264.1	717	No protein	Processed transcript	-	-	TSL:5

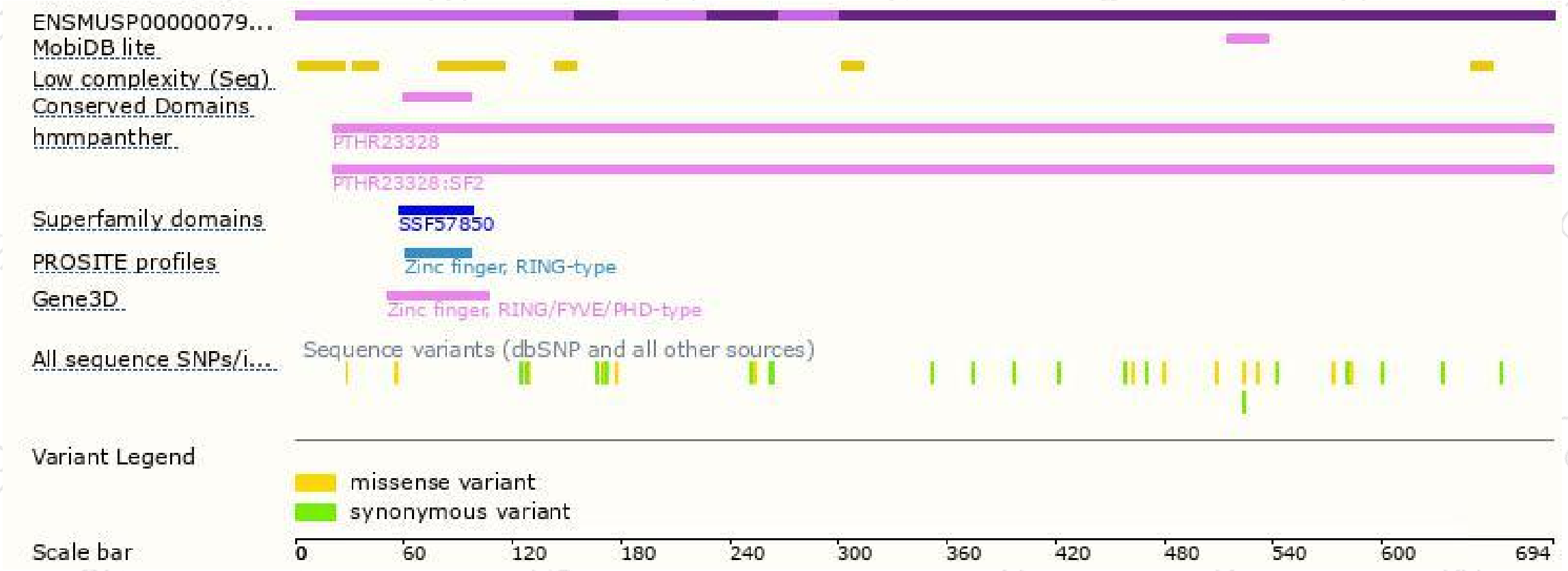
The strategy is based on the design of *Rnf169-201* transcript,The transcription is shown below



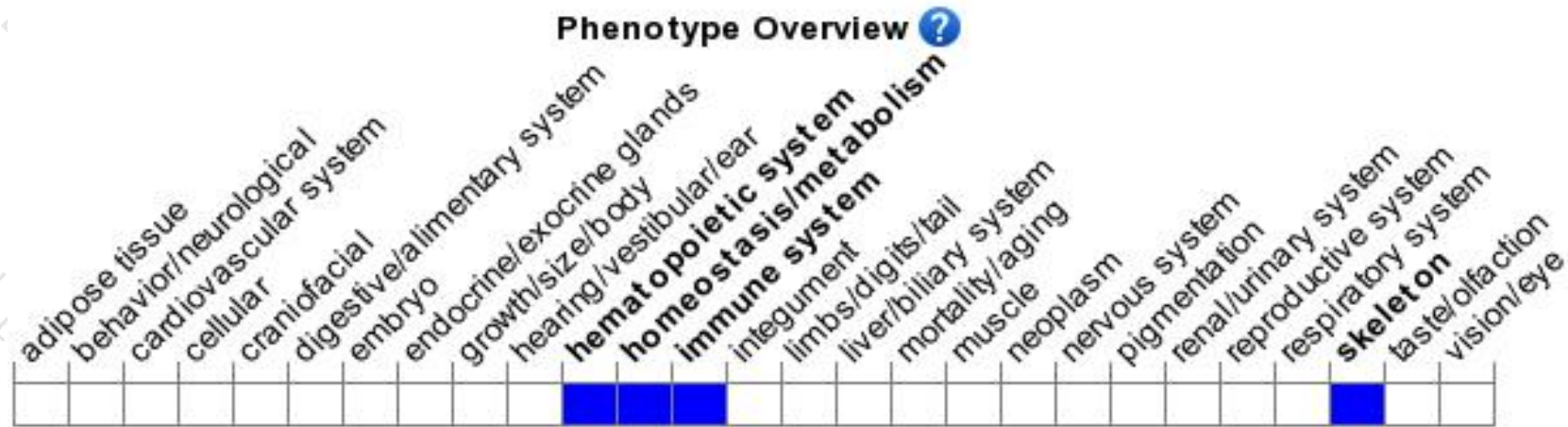
Genomic location distribution



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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