

***Rnf126* Cas9-CKO Strategy**

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Project Overview

Project Name

Rnf126

Project type

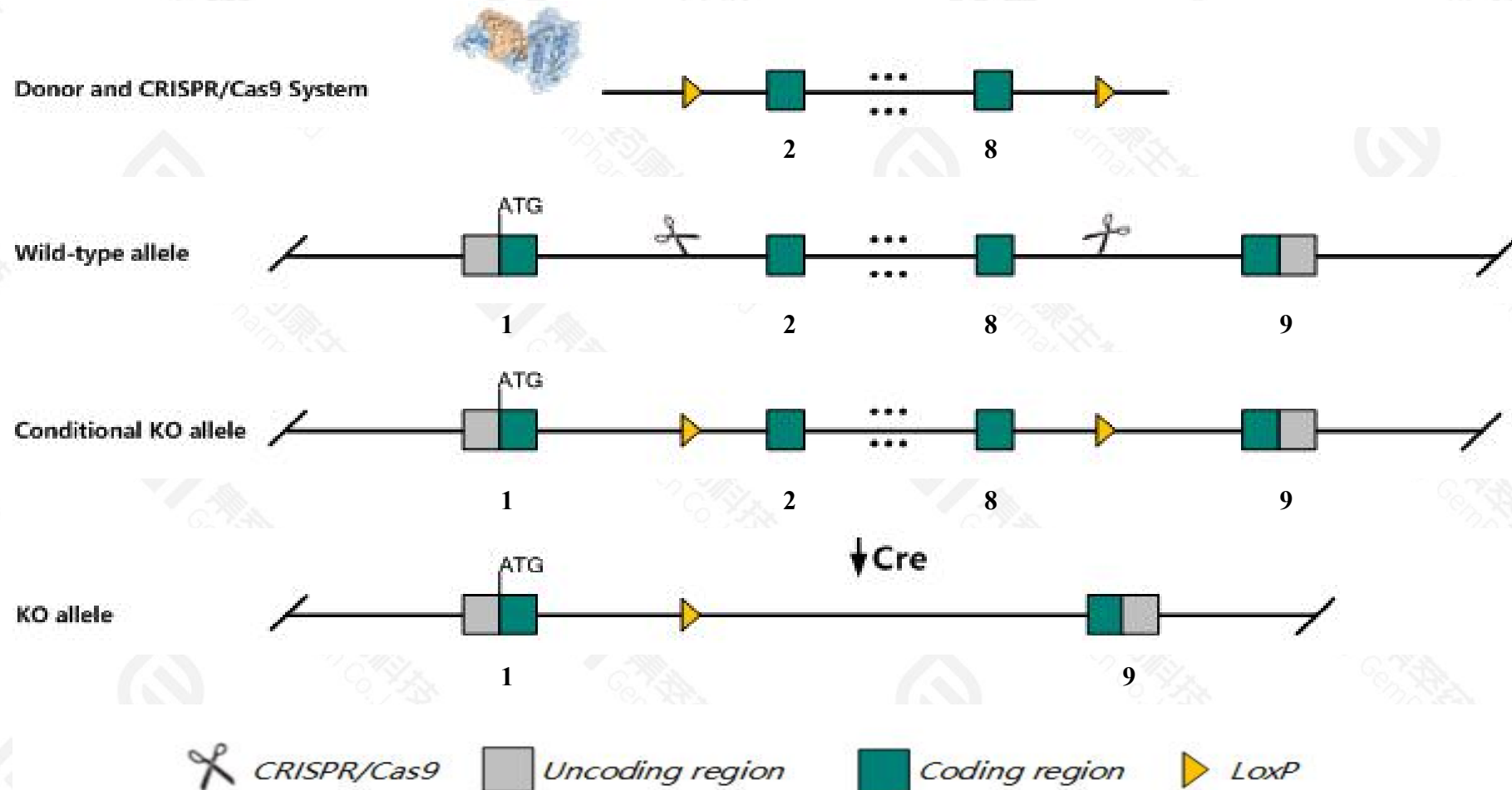
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rnfl26* gene. The schematic diagram is as follows:



- The *Rnf126* gene has 3 transcripts. According to the structure of *Rnf126* gene, exon2-exon8 of *Rnf126-201*(ENSMUST00000047203.9) transcript is recommended as the knockout region. The region contains most of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rnf126* gene. The brief process is as follows: CRISPR/Cas9 system and Donor were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

- The flox mice will not frame-shift mutation after mating with mice expressing Cre recombinase, but most of the coding region was deleted.
- The *Rnf126* gene is located on the Chr10. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Rnf126 ring finger protein 126 [Mus musculus (house mouse)]

Gene ID: 70294, updated on 17-Dec-2020

Summary



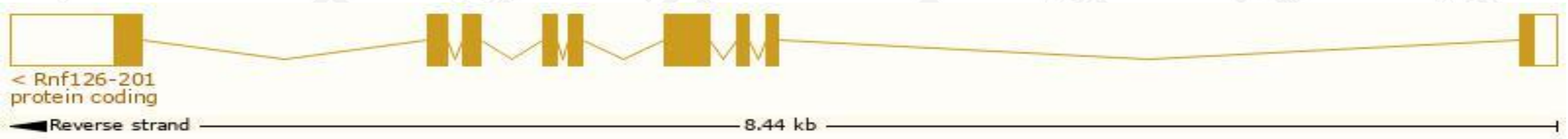
Official Symbol	Rnf126 provided by MGI
Official Full Name	ring finger protein 126 provided by MGI
Primary source	MGI:MGI:1917544
See related	Ensembl:ENSMUSG00000035890
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2610010O19Rik
Expression	Broad expression in testis adult (RPKM 392.3), ovary adult (RPKM 96.9) and 25 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

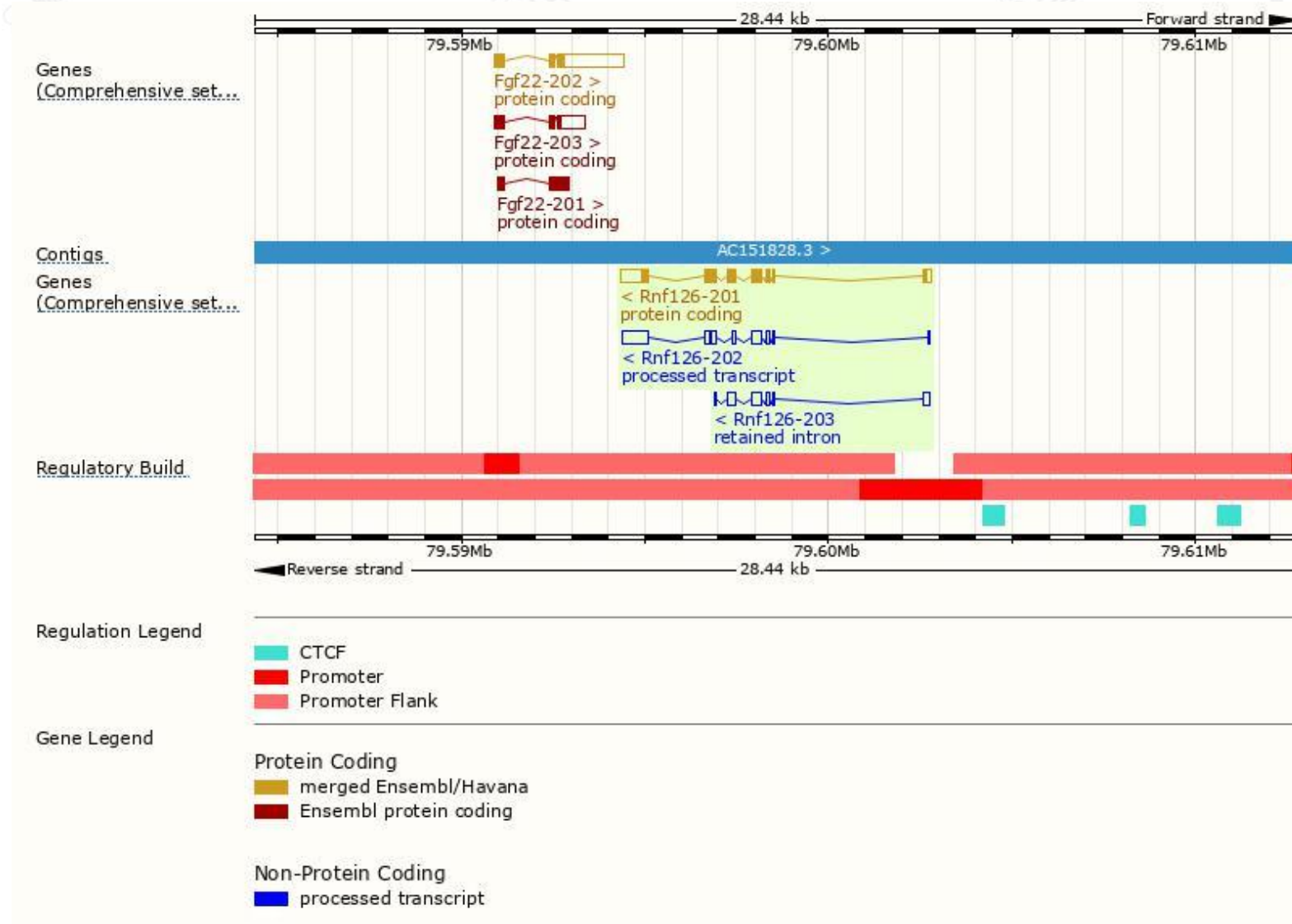
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rnf126-201	ENSMUST00000047203.9	1639	313aa	Protein coding	CCDS23989		TSL:1 , GENCODE basic , APPRIS P1 ,
Rnf126-202	ENSMUST00000218770.2	1408	No protein	Processed transcript	-		TSL:1 ,
Rnf126-203	ENSMUST00000219189.2	773	No protein	Retained intron	-		TSL:3 ,

The strategy is based on the design of *Rnf126-201* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

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