

Rtl6 Cas9-KO Strategy

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Project Overview

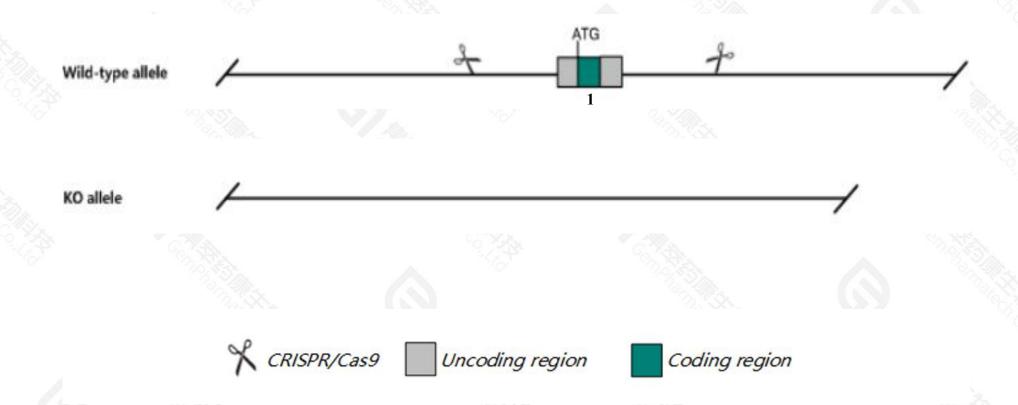


Project Name	Rtl6
Project type	Cas9-KO
Strain background	C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Rtl6* gene. The schematic diagram is as follows:



Technical routes



- > The *Rtl6* gene has 1 transcript. According to the structure of *Rtl6* gene, exon1 of *Rtl6-201*(ENSMUST00000069476.4) transcript is recommended as the knockout region. The region contains all 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 *Rtl6* gene. The brief process is as follows: CRISPR/Cas9 system 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.

Notice



- > The *Rtl6* gene is located on the Chr15. 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)



Rtl6 retrotransposon Gag like 6 [Mus musculus (house mouse)]

Gene ID: 223732, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Rtl6 provided by MGI

Official Full Name retrotransposon Gag like 6 provided by MGI

Primary source MGI:MGI:2675858

See related Ensembl:ENSMUSG00000055745

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 BC058638, Ldoc1l, Mar6, Mart6

Orthologs <u>human</u> all

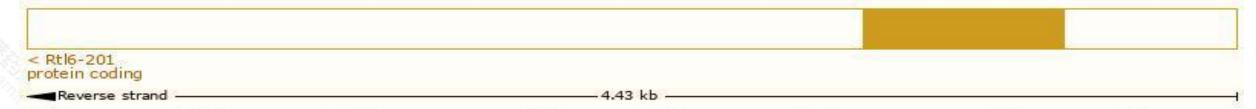
Transcript information (Ensembl)



The gene has 1 transcript, and the transcript is shown below:

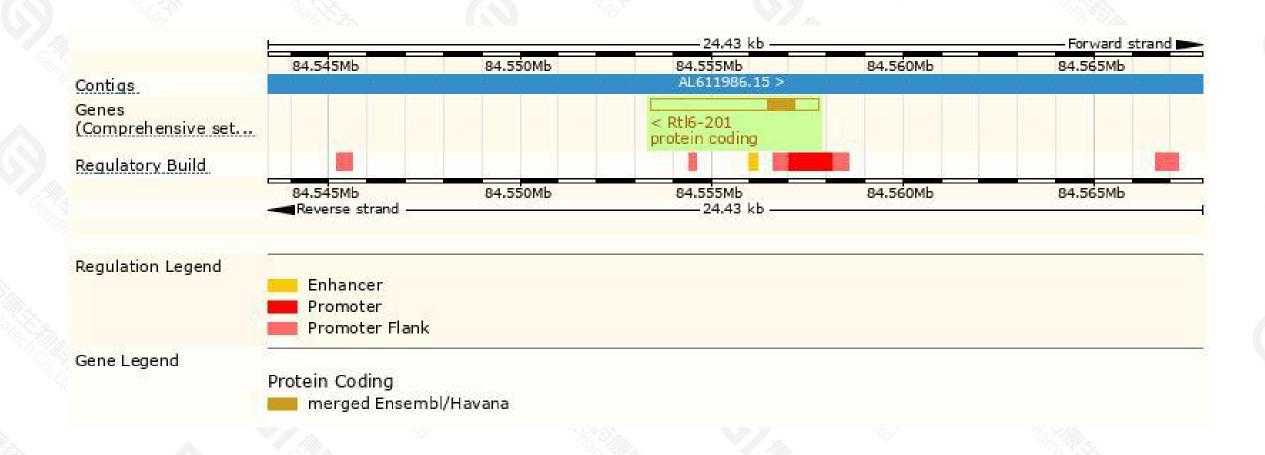
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Rtl6-201	ENSMUST00000069476.4	4426	243aa	Protein coding	CCDS37168	Q505G4	TSL:NA GENCODE basic APPRIS P1	

The strategy is based on the design of *Rtl6-201* transcript, the transcription is shown below:



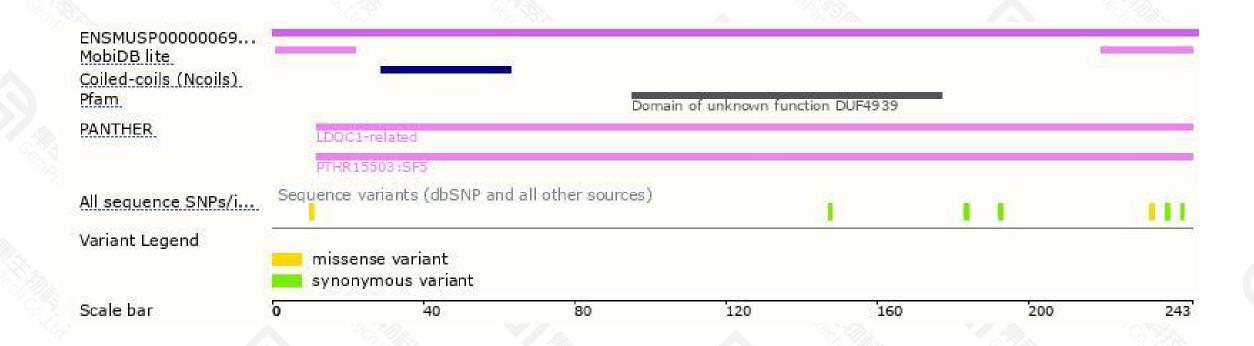
Genomic location distribution





Protein domain







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

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