

Larp6 Cas9-CKO Strategy

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Project Overview



Project Name

Larp6

Project type

Cas9-CKO

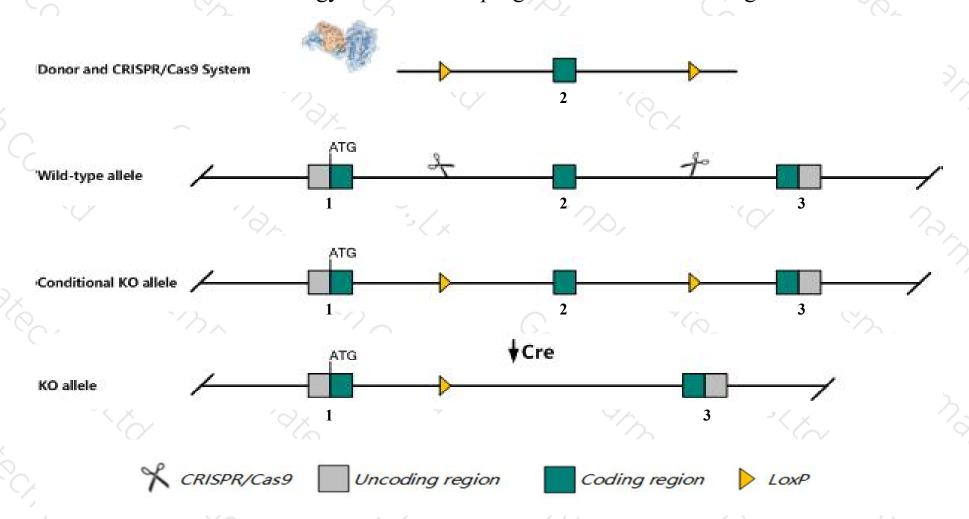
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Larp6 gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Larp6* gene has 1 transcript. According to the structure of *Larp6* gene, exon2 of *Larp6-201*(ENSMUST00000038407.5) transcript is recommended as the knockout region. The region contains 211bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Larp6* 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.

Notice



- > The *Larp6* gene is located on the Chr9. 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)



Larp6 La ribonucleoprotein domain family, member 6 [Mus musculus (house mouse)]

Gene ID: 67557, updated on 31-Jan-2019

Summary

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Official Symbol Larp6 provided by MGI

Official Full Name La ribonucleoprotein domain family, member 6 provided by MGI

Primary source MGI:MGI:1914807

See related Ensembl: ENSMUSG00000034839

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 5430431G03Rik, Al552438, Achn

Expression Broad expression in cerebellum adult (RPKM 12.1), cortex adult (RPKM 9.2) and 16 other tissuesSee more

Orthologs human all

Transcript information (Ensembl)



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

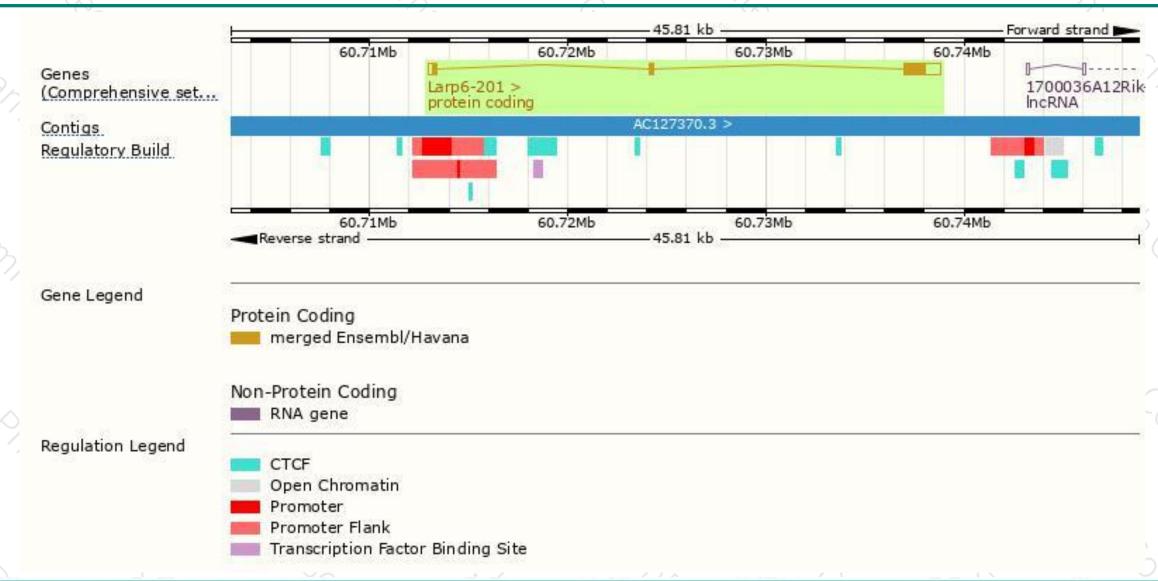
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Larp6-201	ENSMUST00000038407.5	2440	492aa	Protein coding	CCDS23258	Q8BN59	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of Larp6-201 transcript, The transcription is shown below



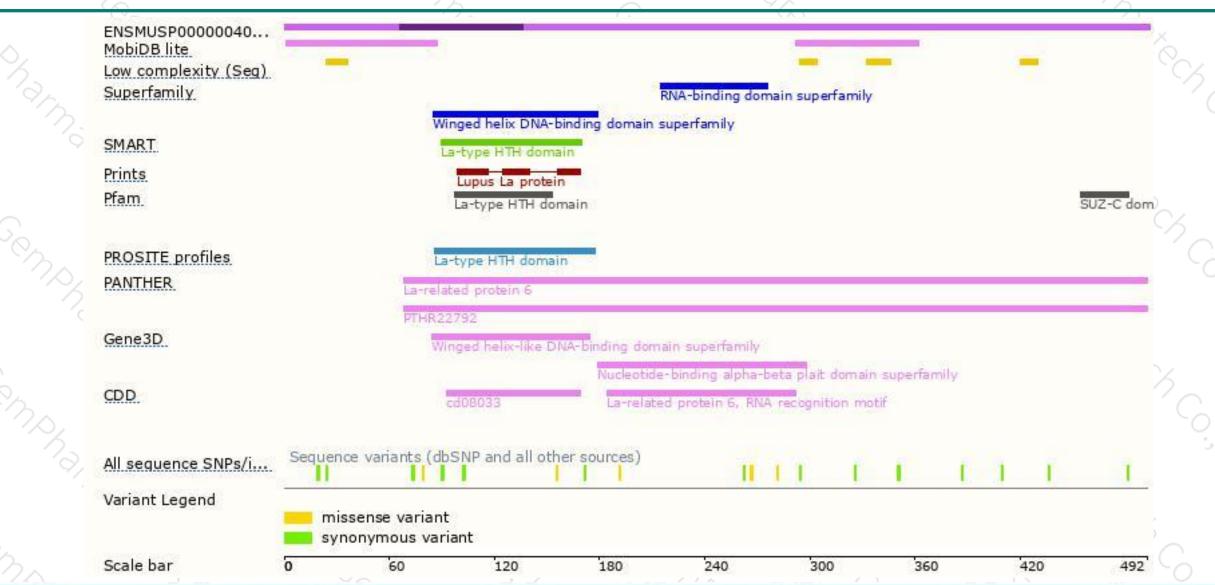
Genomic location distribution





Protein domain







If you have any questions, you are welcome to inquire. Tel: 400-9660890





