

Lcn8 Cas9-KO Strategy

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Project Overview



Project Name

Lcn8

Project type

Cas9-KO

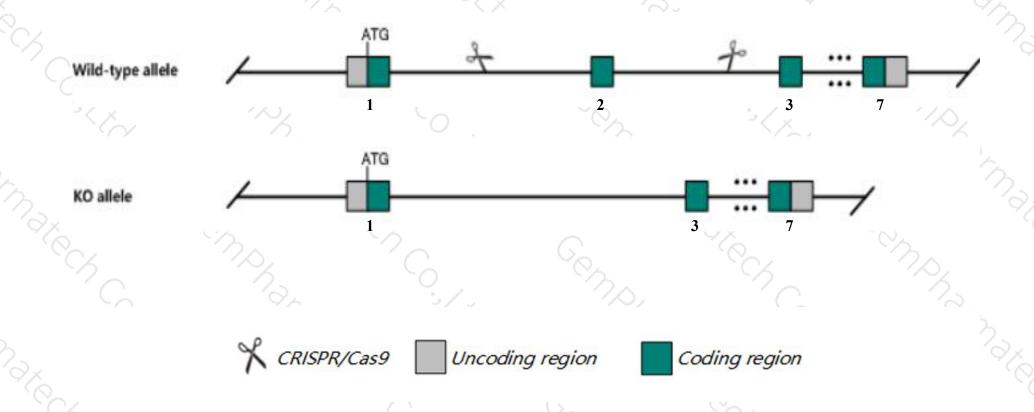
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Lcn8* gene. The schematic diagram is as follows:



Technical routes



- > The *Lcn8* gene has 1 transcript. According to the structure of *Lcn8* gene, exon2 of *Lcn8-201*(ENSMUST00000038482.6) transcript is recommended as the knockout region. The region contains 131bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Lcn8* 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 Intron2 is only 453bp,loxp insertion may affect mRNA splicing.
- \succ The KO region contains functional region of the *Lcn5* gene. Knockout the region may affect the function of *Lcn5* gene.
- > The *Lcn8* 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)



Lcn8 lipocalin 8 [Mus musculus (house mouse)]

Gene ID: 78076, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Lcn8 provided by MGI

Official Full Name lipocalin 8 provided by MGI

Primary source MGI:MGI:2135945

See related Ensembl: ENSMUSG00000036449

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 9230106L18Rik, EP17, Lcn5, mEP17

Expression Restricted expression toward genital fat pad adult (RPKM 2151.2)See more

Orthologs <u>human all</u>

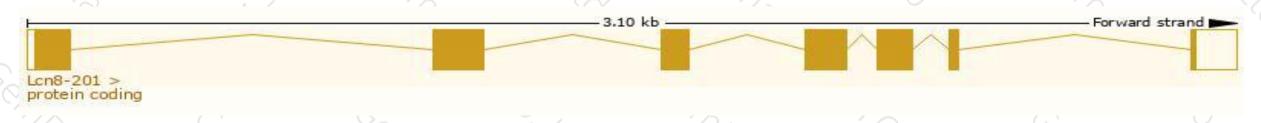
Transcript information (Ensembl)



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

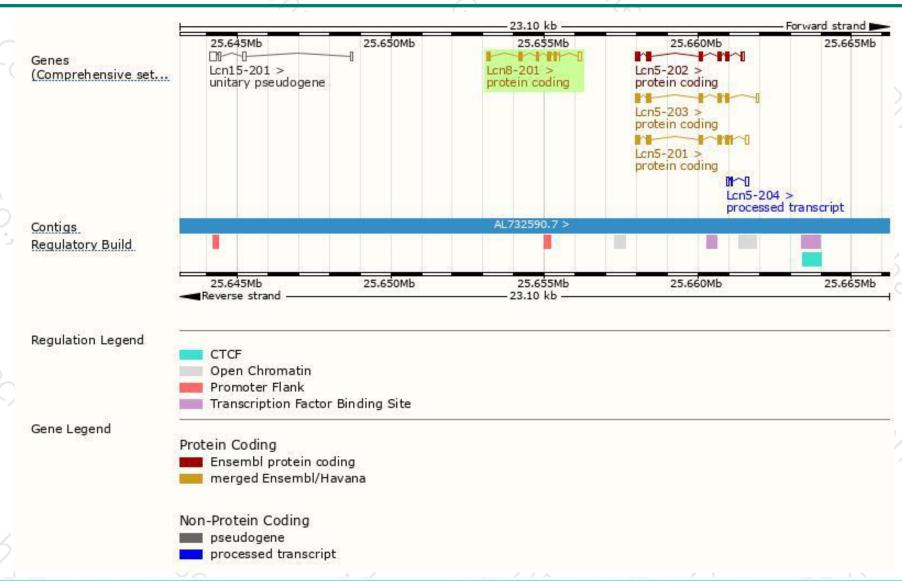
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Lcn8-201	ENSMUST00000038482.6	651	<u>175aa</u>	Protein coding	CCDS15786	Q924P3	TSL:1 GENCODE basic APPRIS P1	

The strategy is based on the design of *Lcn8-201* transcript, the transcription is shown below:



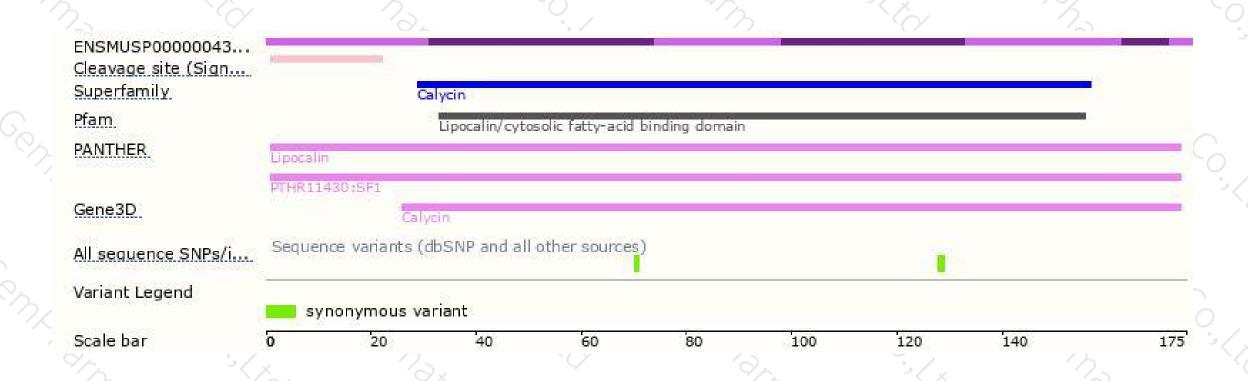
Genomic location distribution





Protein domain







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





