

Kdelr2 Cas9-CKO Strategy

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Design Date: 2019/9/11

Project Overview



Project Name

Kdelr2

Project type

Cas9-CKO

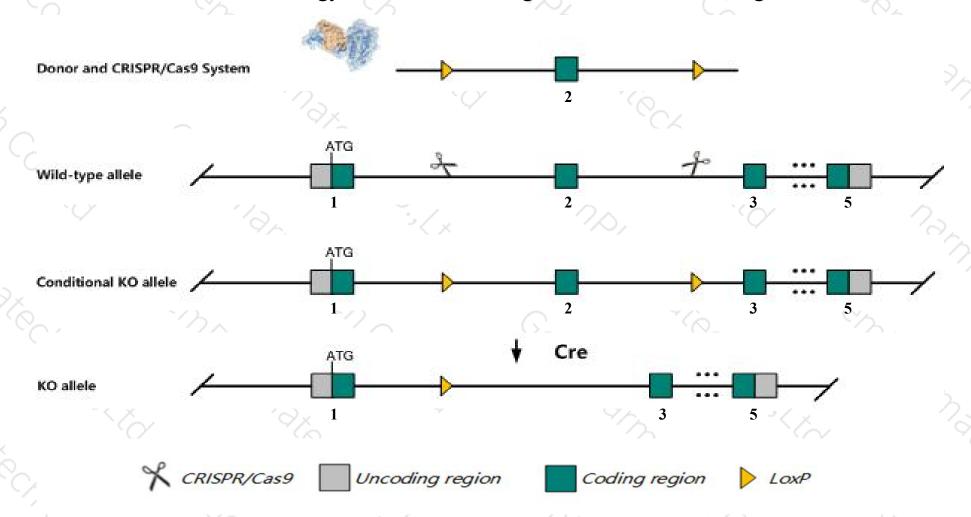
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Kdelr2* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Kdelr2* gene has 1 transcript. According to the structure of *Kdelr2* gene, exon2 of *Kdelr2-201*(ENSMUST00000110731.3) transcript is recommended as the knockout region. The region contains 101bp coding sequence.

 Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Kdelr2* 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 *Kdelr2* gene is located on the Chr5. 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)



Kdelr2 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 [Mus musculus (house mouse)]

Gene ID: 66913, updated on 14-Aug-2019

Summary

△ ?

Official Symbol Kdelr2 provided by MGI

Official Full Name KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 provided by MGI

Primary source MGI:MGI:1914163

See related Ensembl: ENSMUSG00000079111

Gene type protein coding
RefSeq status VALIDATED
Organism <u>Mus musculus</u>

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as 1110007A14Rik

Expression Ubiquitous expression in duodenum adult (RPKM 116.2), colon adult (RPKM 108.7) and 28 other tissues See more

Orthologs <u>human</u> all

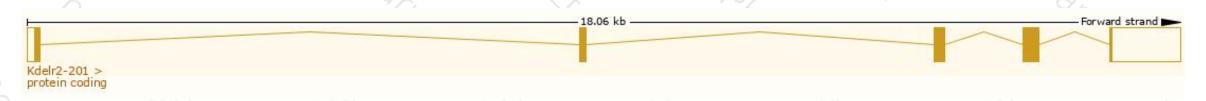
Transcript information (Ensembl)



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

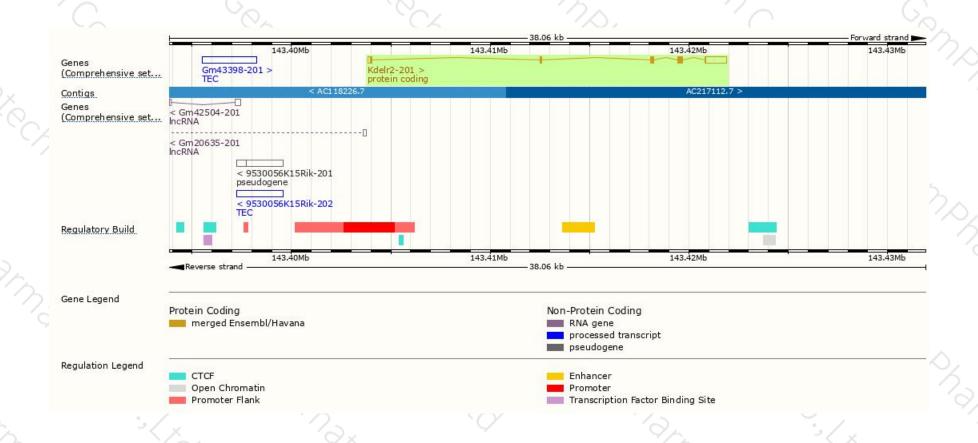
Name 🍦	Transcript ID	bp 🍦	Protein	Biotype	CCDS	UniProt 🍦	Flags		
Kdelr2-201	ENSMUST00000110731.3	1829	212aa	Protein coding	CCDS39369₽	Q9CQM2₺	TSL:1	GENCODE basic	APPRIS P1

The strategy is based on the design of *Kdelr2-201* transcript, The transcription is shown below



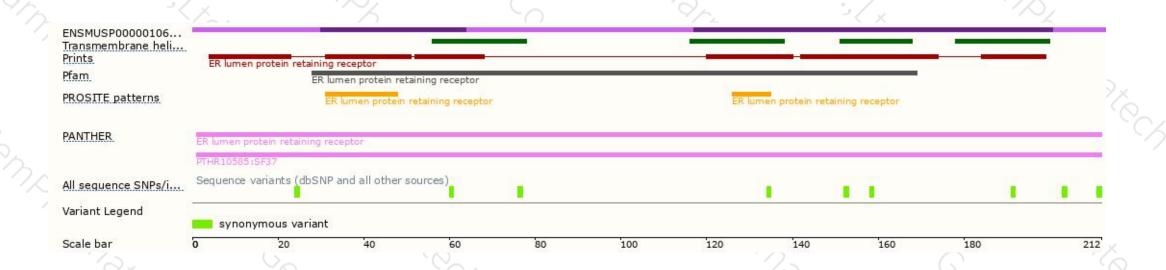
Genomic location distribution





Protein domain







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





