

***Kdelr3* Cas9-CKO Strategy**

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Project Overview

Project Name

Kdelr3

Project type

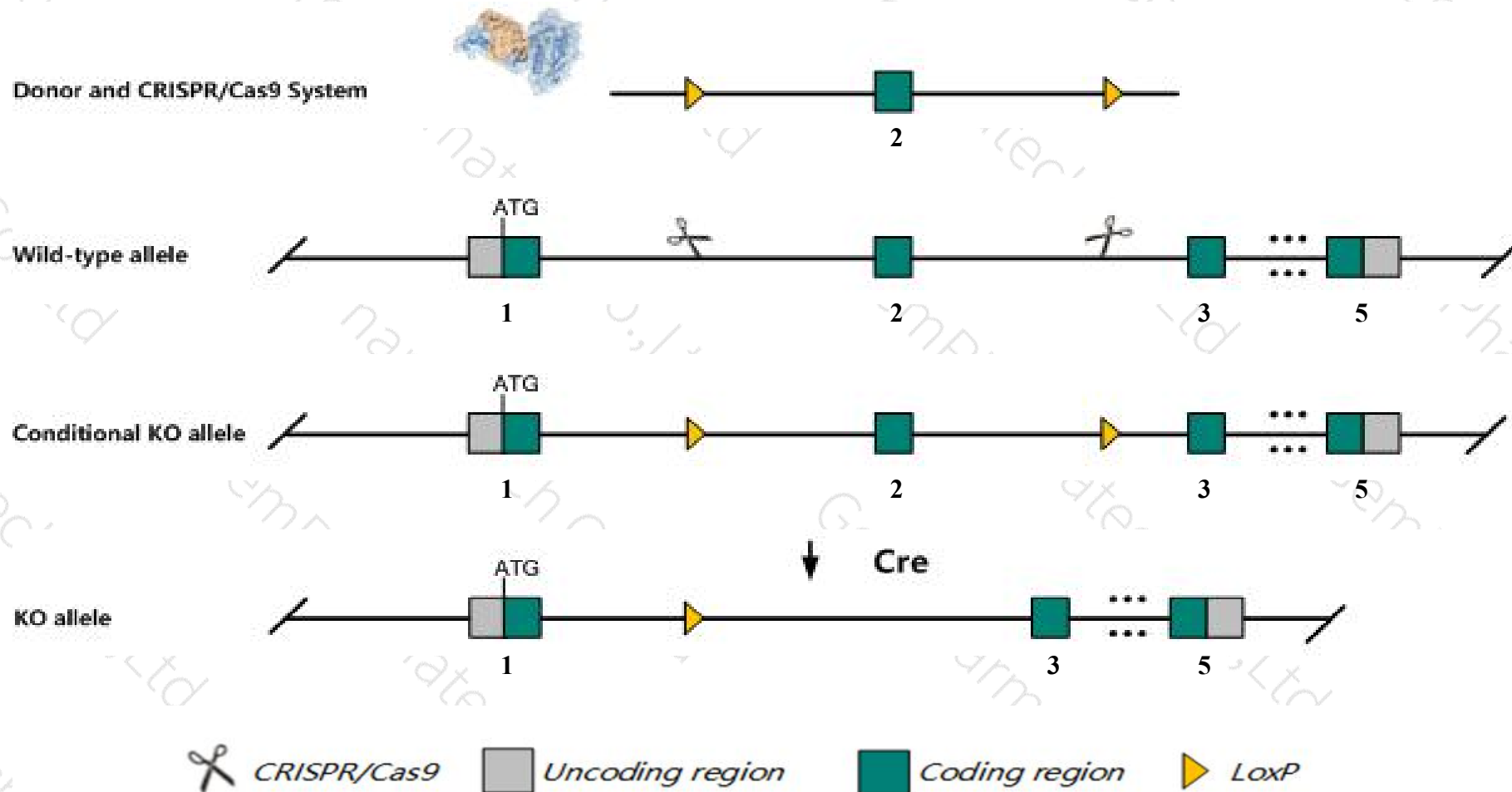
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Kdelr3* gene. The schematic diagram is as follows:



- The *Kdelr3* gene has 1 transcript. According to the structure of *Kdelr3* gene, exon2 of *Kdelr3-201* (ENSMUST00000010974.8) 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 *Kdelr3* 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 *Kdelr3* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Kdelr3 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3 [Mus musculus (house mouse)]

Gene ID: 105785, updated on 13-Mar-2020

Summary



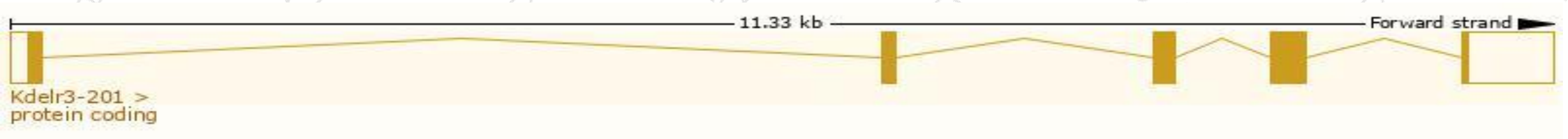
Official Symbol	Kdelr3 provided by MGI
Official Full Name	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3 provided by MGI
Primary source	MGI:MGI:2145953
See related	Ensembl:ENSMUSG00000010830
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	A1173274
Expression	Broad expression in limb E14.5 (RPKM 49.3), subcutaneous fat pad adult (RPKM 21.0) and 17 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

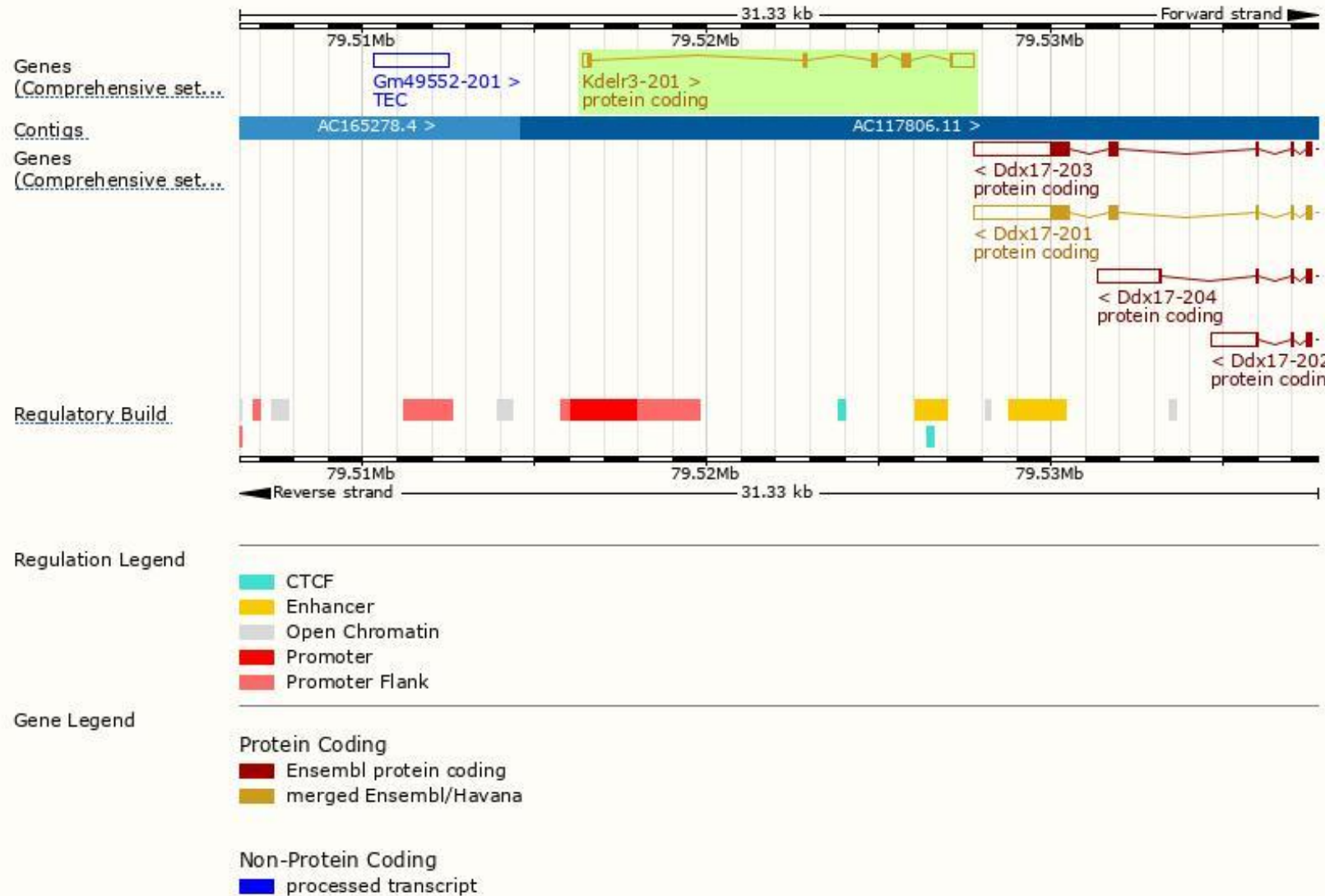
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Kdelr3-201	ENSMUST00000010974.8	1411	214aa	Protein coding	CCDS27642	Q8R1L4	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Kdelr3-201* 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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