

Dmac1 Cas9-CKO Strategy

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Project Overview



Project Name

Dmac1

Project type

Cas9-CKO

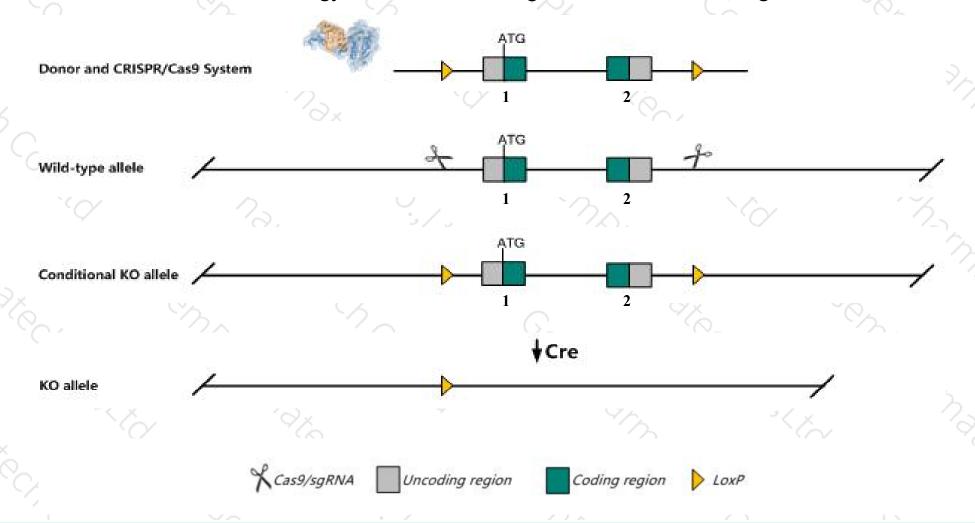
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Dmac1* gene. The schematic diagram is as follows:



Technical routes



- > The *Dmac1* gene has 1 transcript. According to the structure of *Dmac1* gene, exon1-exon2 of *Dmac1*201(ENSMUST00000030103.8) 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 *Dmac1* gene. The brief process is as follows:sgRNA was transcribed in vitro, donor vector was constructed.Cas9, sgRNA 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 was 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.

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Notice



- > The *Dmac1* gene is located on the Chr4. 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)



Dmac1 distal membrane arm assembly complex 1 [Mus musculus (house mouse)]

Gene ID: 66928, updated on 25-Sep-2020

Summary

↑ ?

Official Symbol Dmac1 provided by MGI

Official Full Name distal membrane arm assembly complex 1 provided by MGI

Primary source MGI:MGI:1914178

See related Ensembl: ENSMUSG00000028398

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 Tmem261; 1700027K24Rik; 3110001D03Rik

Expression Ubiquitous expression in thymus adult (RPKM 28.4), limb E14.5 (RPKM 27.6) and 28 other tissues See more

Orthologs human all

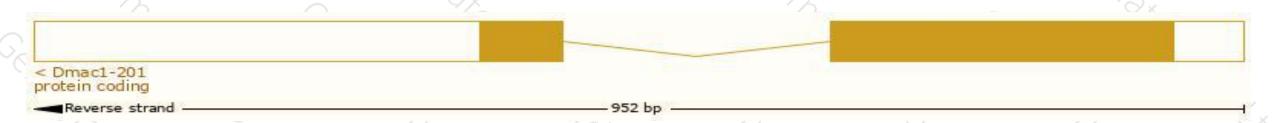
Transcript information (Ensembl)



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

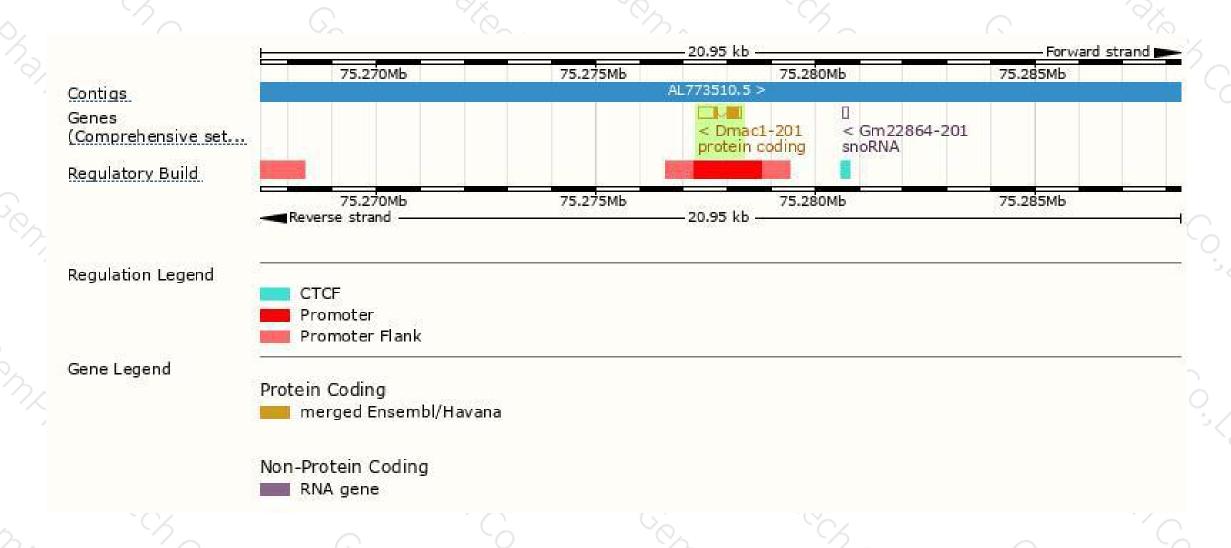
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Dmac1-201	ENSMUST00000030103.8	741	<u>111aa</u>	Protein coding	CCDS18288	Q9CQ00	TSL:1 GENCODE basic APPRIS P1	L

The strategy is based on the design of *Dmac1-201* transcript, the transcription is shown below:



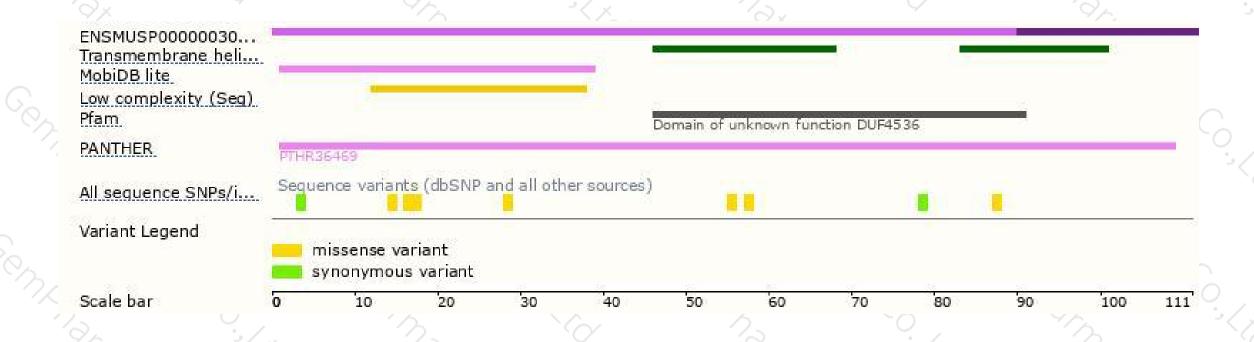
Genomic location distribution





Protein domain







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

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