

Carnmt1 Cas9-CKO Strategy

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Design Date: 2020-5-28

Project Overview



Project Name Carnmt1

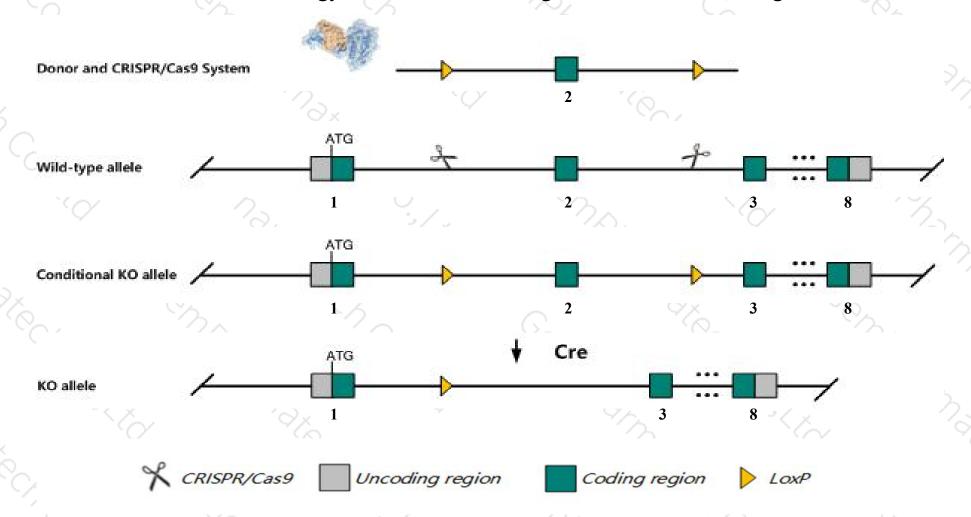
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Carnmt1 gene. The schematic diagram is as follows:



Technical routes



- The *Carnmt1* gene has 3 transcripts. According to the structure of *Carnmt1* gene, exon2 of *Carnmt1-201* (ENSMUST00000025632.10) transcript is recommended as the knockout region. The region contains 196bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Carnmt1* 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 *Carnmt1* gene is located on the Chr19. 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)



Carnmt1 carnosine N-methyltransferase 1 [Mus musculus (house mouse)]

Gene ID: 67383, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Carnmt1 provided by MGI

Official Full Name carnosine N-methyltransferase 1 provided by MGI

Primary source MGI:MGI:1914633

See related Ensembl: ENSMUSG00000024726

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 2410127L17Rik

Expression Ubiquitous expression in ovary adult (RPKM 5.9), liver E14 (RPKM 5.1) and 28 other tissuesSee more

Orthologs <u>human</u> all

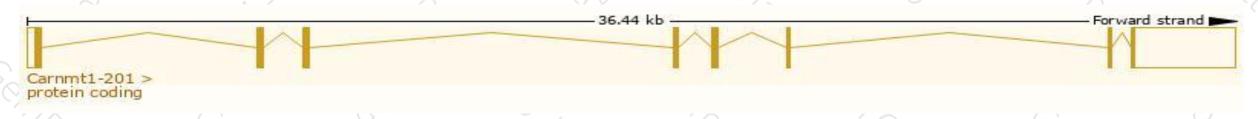
Transcript information (Ensembl)



The gene has 3 transcripts, all transcripts are shown below:

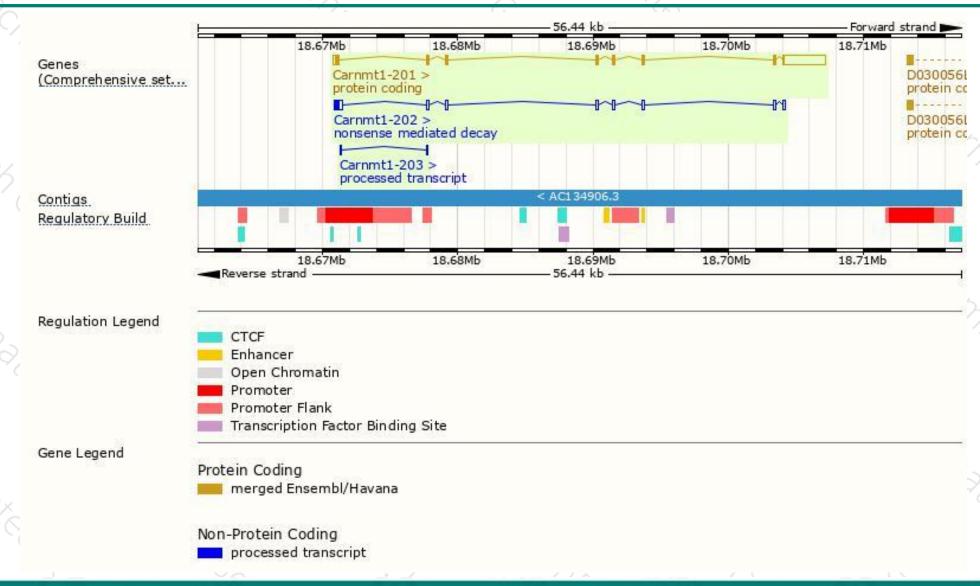
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Carnmt1-201	ENSMUST00000025632.10	4518	400aa	Protein coding	CCDS29689	Q80UY1	TSL:1 GENCODE basic APPRIS P1
Carnmt1-202	ENSMUST00000235736.1	1714	<u>68aa</u>	Nonsense mediated decay	-	A0A494BA54	
Carnmt1-203	ENSMUST00000236401.1	215	No protein	Processed transcript	12	ų.	

The strategy is based on the design of Carnmt1-201 transcript, the transcription is shown below:



Genomic location distribution





Protein domain







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





