

Camk4 Cas9-CKO Strategy

Designer: Daohua Xu

Reviewer: Huimin Su

Design Date: 2019-11-22

Project Overview



Project Name Camk4

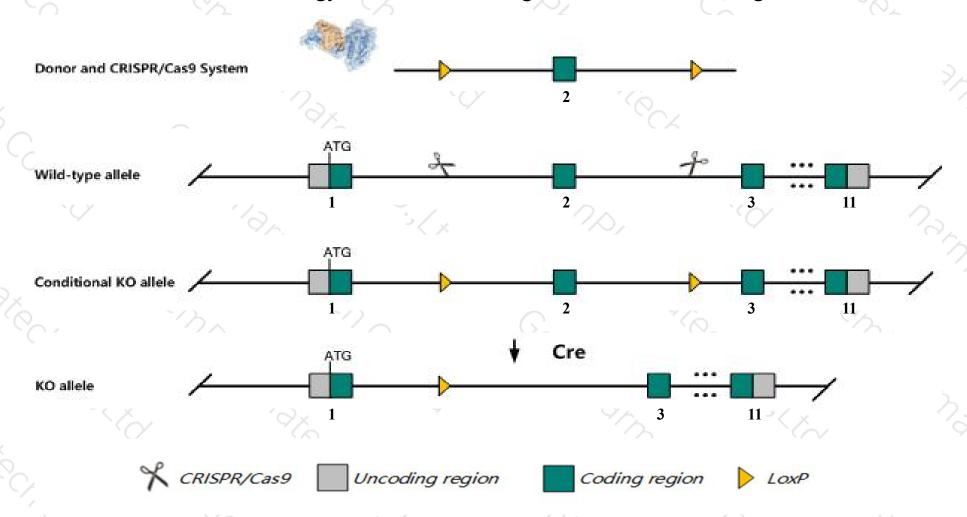
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Camk4 gene. The schematic diagram is as follows:



Technical routes



- The Camk4 gene has 4 transcripts. According to the structure of Camk4 gene, exon2 of Camk4-201 (ENSMUST00000042868.5) transcript is recommended as the knockout region. The region contains 79bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Camk4* 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



- > According to the existing MGI data, Homozygotes for different targeted mutations show variable phenotypes, including reduced viability, male and/or female sterility, and mild to severe neurological and spatial memory disorders.
- > The Camk4 gene is located on the Chr18. 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)



Camk4 calcium/calmodulin-dependent protein kinase IV [Mus musculus (house mouse)]

Gene ID: 12326, updated on 5-Mar-2019

Summary

☆ ?

Official Symbol Camk4 provided by MGI

Official Full Name calcium/calmodulin-dependent protein kinase IV provided by MGI

Primary source MGI:MGI:88258

See related Ensembl:ENSMUSG00000038128

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 A430110E23Rik, Al666733, CaMKIV, CaMKIV/Gr, D18Bwg0362e

Expression Biased expression in cerebellum adult (RPKM 21.8), cortex adult (RPKM 9.3) and 6 other tissuesSee more

Orthologs <u>human</u> all

Transcript information (Ensembl)



The gene has 4 transcripts, all transcripts are shown below:

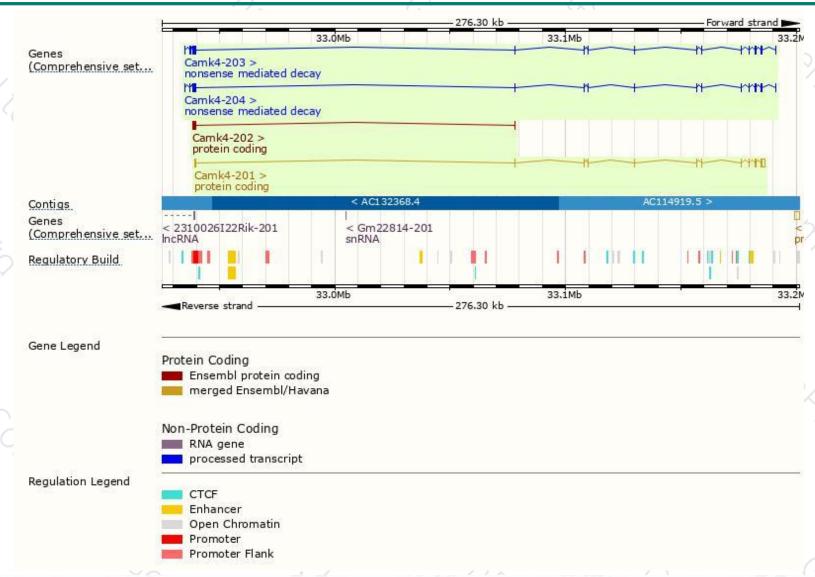
		h,					
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Camk4-201	ENSMUST00000042868.5	2717	469aa	Protein coding	CCDS29122	Q8BGR3	TSL:1 GENCODE basic APPRIS P1
Camk4-202	ENSMUST00000234001.1	526	<u>67aa</u>	Protein coding	-	(*)	CDS 3' incomplete
Camk4-203	ENSMUST00000234205.1	1868	469aa	Nonsense mediated decay	CCDS29122	120	
Camk4-204	ENSMUST00000234654.1	1829	469aa	Nonsense mediated decay	CCDS29122	100	
~ / /		V 1997 A				and the same of th	

The strategy is based on the design of Camk4-201 transcript, The transcription is shown below



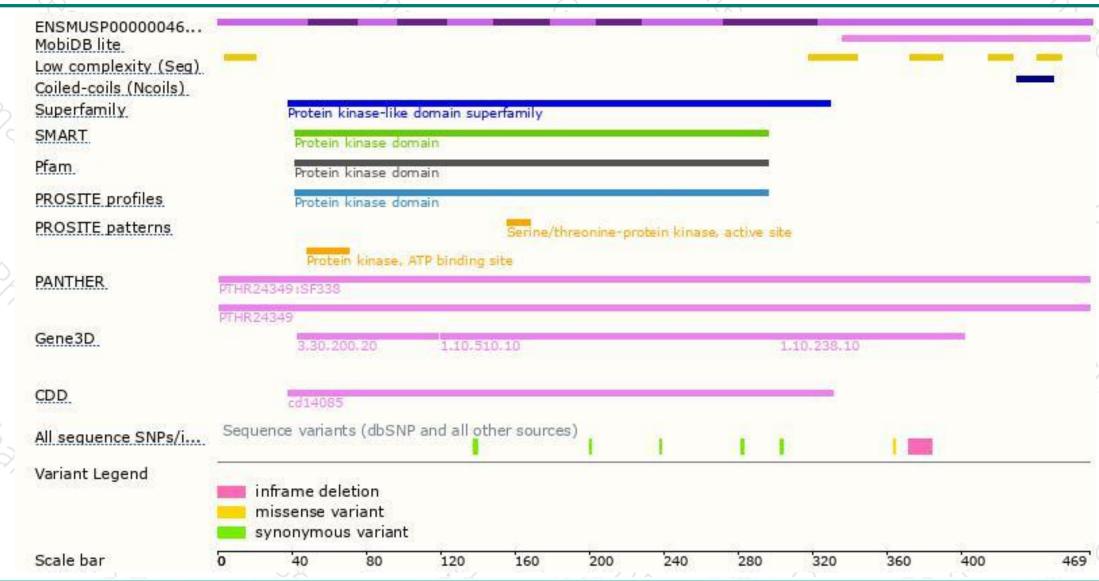
Genomic location distribution





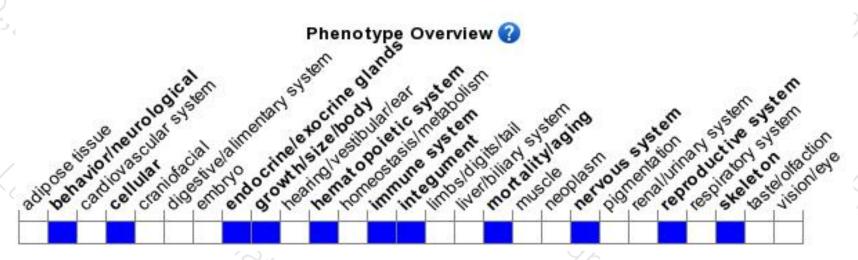
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

According to the existing MGI data, Homozygotes for different targeted mutations show variable phenotypes, including reduced viability, male and/or female sterility, and mild to severe neurological and spatial memory disorders.



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





