

Cdkl1 Cas9-KO Strategy

Designer: Xiaojing Li

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Reviewer:JiaYu

Project Overview



Project Name Cdk11

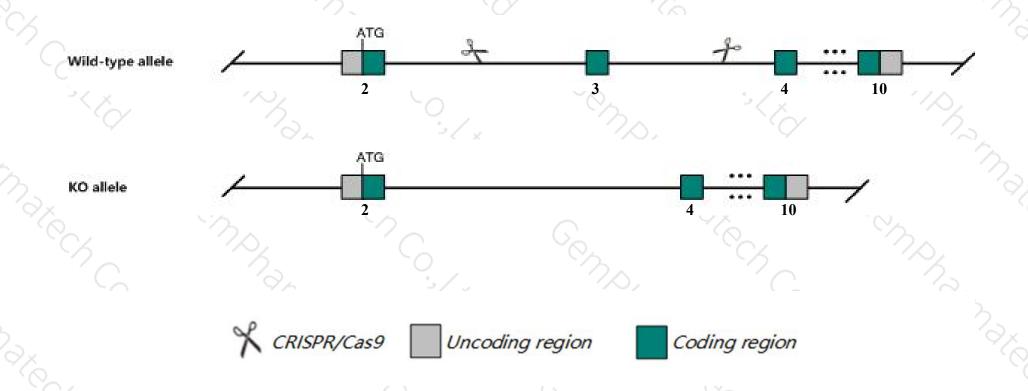
Project type Cas9-KO

Strain background C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Cdkl1* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Cdkl1* gene has 2 transcripts. According to the structure of *Cdkl1* gene, exon3 of *Cdkl1-201*(ENSMUST00000021377.4) transcript is recommended as the knockout region. The region contains 122bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Cdkl1* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- ➤ The *Cdkl1* gene is located on the Chr12. 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)



Cdkl1 cyclin-dependent kinase-like 1 (CDC2-related kinase) [Mus musculus (house mouse)]

Gene ID: 71091, updated on 14-Aug-2019

Summary

Official Symbol Cdkl1 provided by MGI

Official Full Name cyclin-dependent kinase-like 1 (CDC2-related kinase) provided by MGI

Primary source MGI:MGI:1918341

See related Ensembl: ENSMUSG00000020990

RefSeq status PROVISIONAL
Organism Mus musculus

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

Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Also known as 4933411017Rik

Expression Biased expression in kidney adult (RPKM 23.9), ovary adult (RPKM 3.4) and 7 other tissues See more

Orthologs human all

Transcript information (Ensembl)



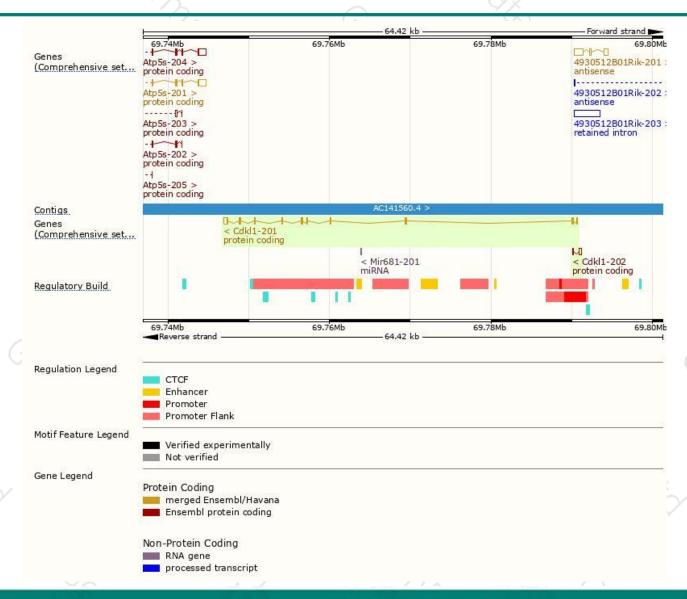
The gene has 2 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cdkl1-201	ENSMUST00000021377.4	1689	352aa	Protein coding	CCDS25955	Q14BG3 Q8CEQ0	TSL:1 GENCODE basic APPRIS P1
Cdkl1-202	ENSMUST00000221646.1	465	<u>1aa</u>	Protein coding	. 5	8-	CDS 3' incomplete TSL:2

The strategy is based on the design of Cdkl1-201 transcript, The transcription is shown below

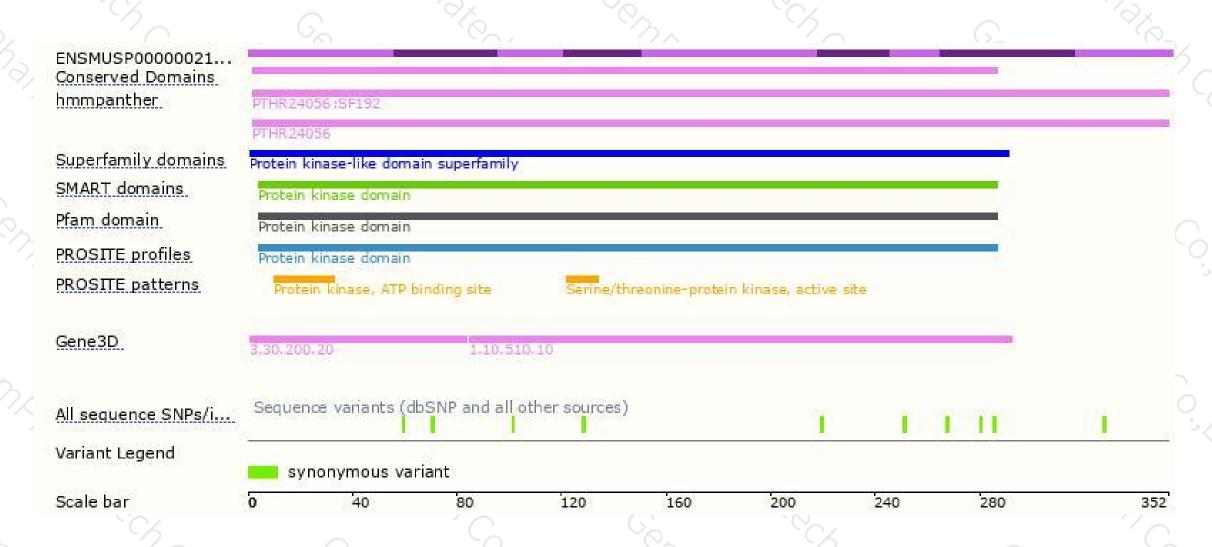
Genomic location distribution





Protein domain







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





