

# Ccnb1ip1 Cas9-CKO Strategy

Designer: Xiaojing Li

Design Date:2019-11-6

Reviewer:JiaYu

## **Project Overview**



**Project Name** 

Ccnb1ip1

**Project type** 

Cas9-CKO

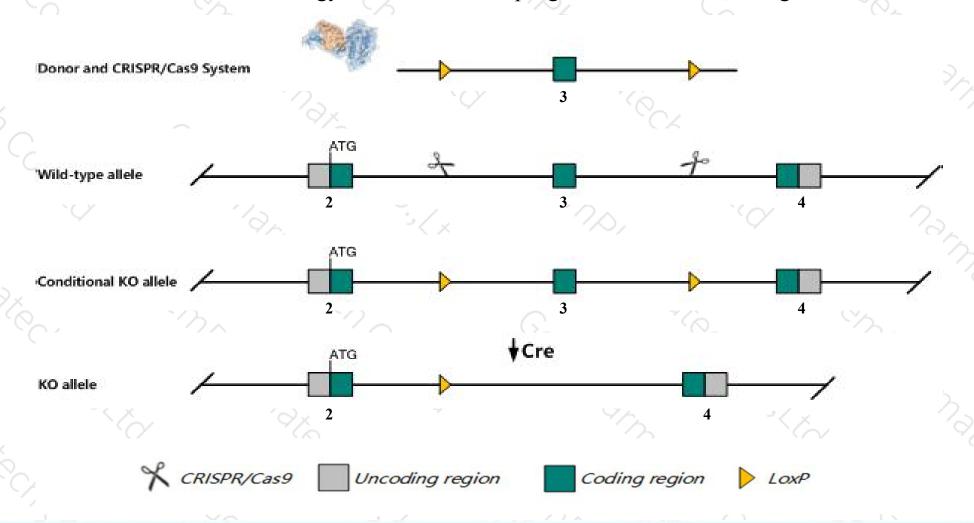
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Ccnblip1* gene. The schematic diagram is as follows:



### Technical routes



- ➤ The *Ccnb1ip1* gene has 2 transcripts. According to the structure of *Ccnb1ip1* gene, exon3 of *Ccnb1ip1-201*(ENSMUST00000095932.4) transcript is recommended as the knockout region. The region contains 334bp coding sequence.

  Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ccnb1ip1* 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, Mice homozygous for an ENU-induced mutation have abnormal testicular and ovarian morphology and exhibit sterility in both sexes owing to meiotic defects.
- The *Ccnblip1* gene is located on the Chr14. 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)



#### Ccnb1ip1 cyclin B1 interacting protein 1 [ Mus musculus (house mouse) ]

Gene ID: 239083, updated on 16-Sep-2019

#### Summary

Official Symbol Ccnb1ip1 provided by MGI

Official Full Name cyclin B1 interacting protein 1 provided by MGI

Primary source MGI:MGI:2685134

See related Ensembl: ENSMUSG00000071470

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 mei4; Gm288; Hei10

Expression Restricted expression toward testis adult (RPKM 2.9) See more

Orthologs human all

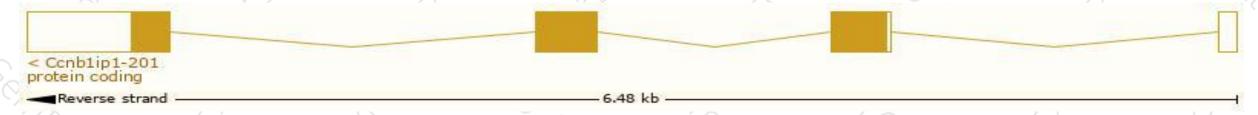
## Transcript information (Ensembl)



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

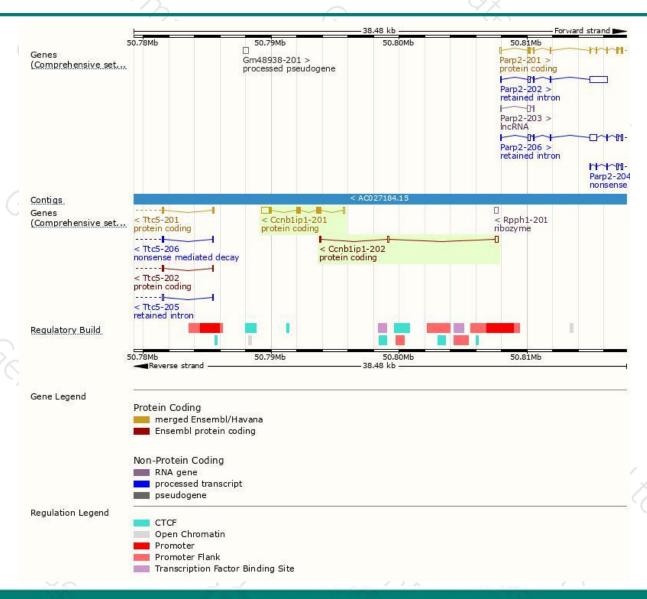
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ccnb1ip1-201	ENSMUST00000095932.4	1509	276aa	Protein coding	CCDS49477	D3Z3K2	TSL:5 GENCODE basic APPRIS P1
Ccnb1ip1-202	ENSMUST00000227614.1	444	<u>16aa</u>	Protein coding		A0A2I3BPZ8	CDS 3' incomplete

The strategy is based on the design of Ccnb1ip1-201 transcript, The transcription is shown below



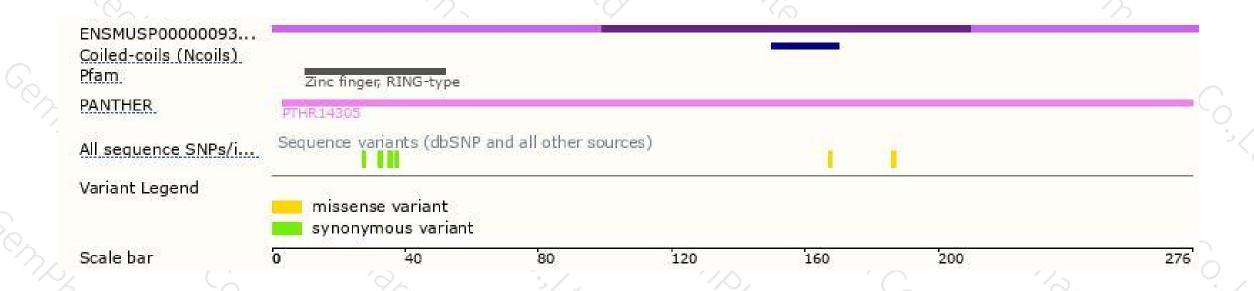
### Genomic location distribution





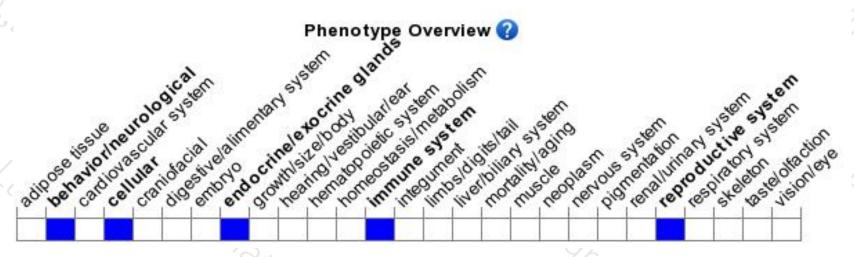
## Protein domain





## 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, Mice homozygous for an ENU-induced mutation have abnormal testicular and ovarian morphology and exhibit sterility in both sexes owing to meiotic defects.



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





