

Ccnblip1 Cas9-KO Strategy

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Project Overview

Project Name

Ccnblip1

Project type

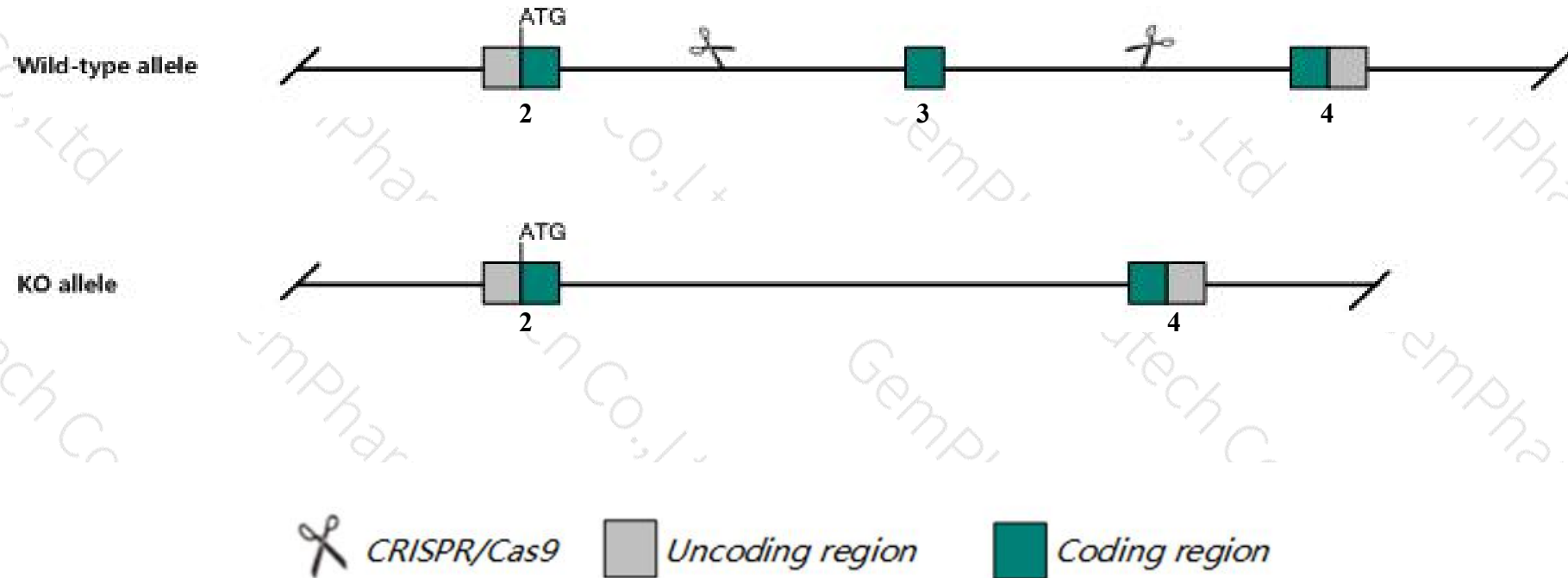
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- The *Ccnblip1* gene has 2 transcripts. According to the structure of *Ccnblip1* gene, exon3 of *Ccnblip1-201* (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 *Ccnblip1* gene. The brief process is as follows: CRISPR/Cas9 system

- 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

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

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

Summary

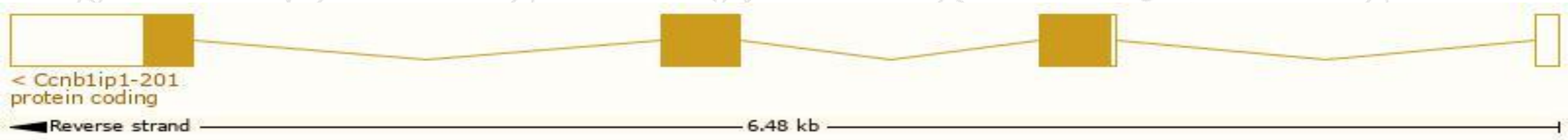
Official Symbol	Ccnb1ip1 provided by MGI
Official Full Name	cyclin B1 interacting protein 1 provided by MGI
Primary source	MGI:MGI:2685134
See related	Ensembl:ENSMUSG000000071470
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; 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)

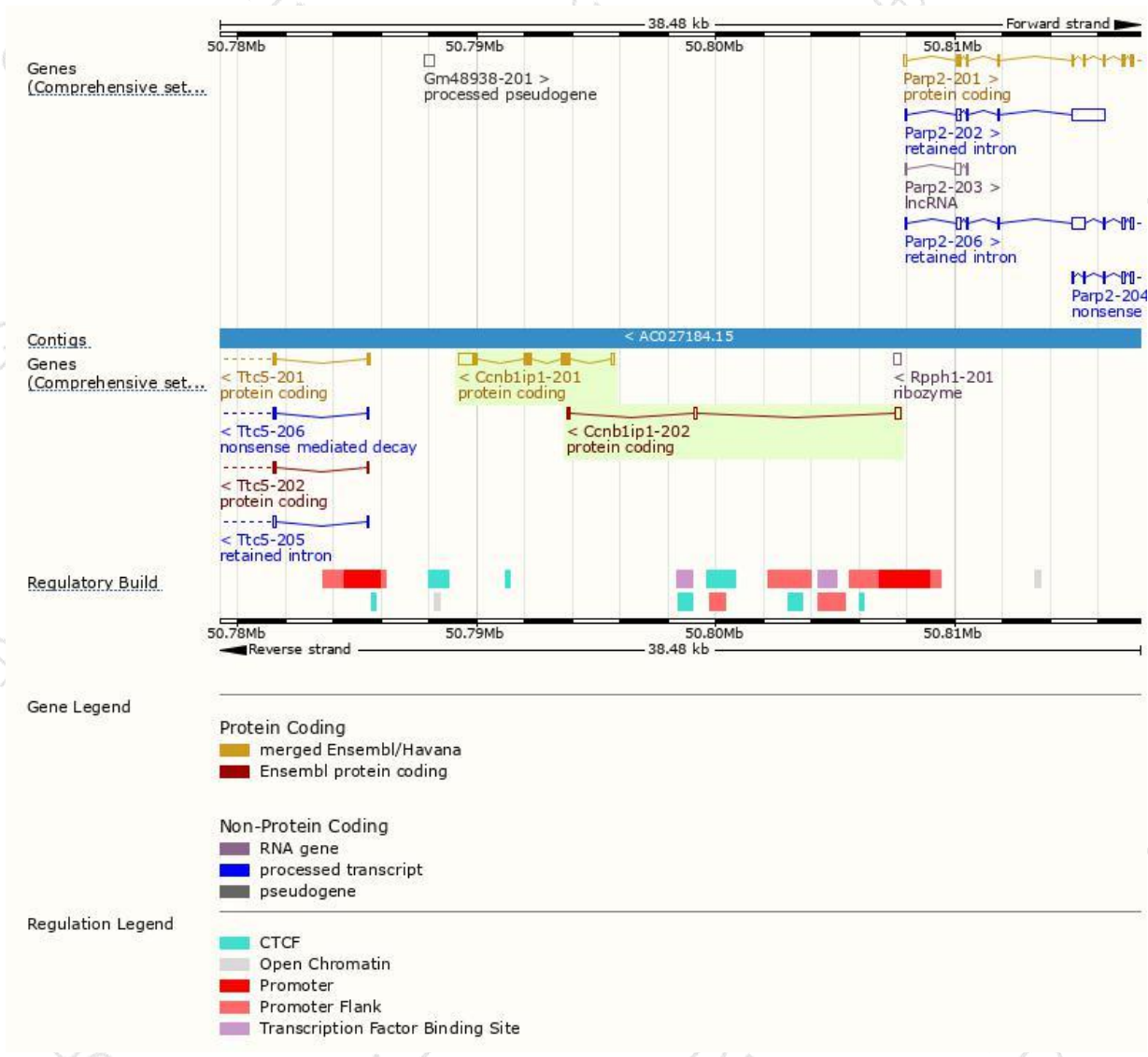
The gene has 2 transcripts,all transcripts are shown below:

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	16aa	Protein coding	-	A0A2I3BPZ8	CDS 3' incomplete

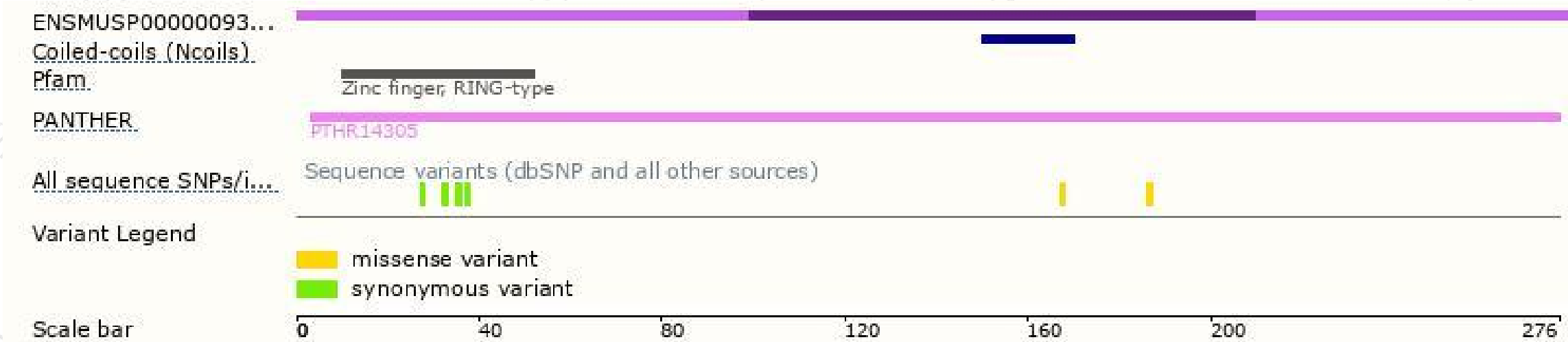
The strategy is based on the design of *Ccnb1ip1-201* transcript,The transcription is shown below



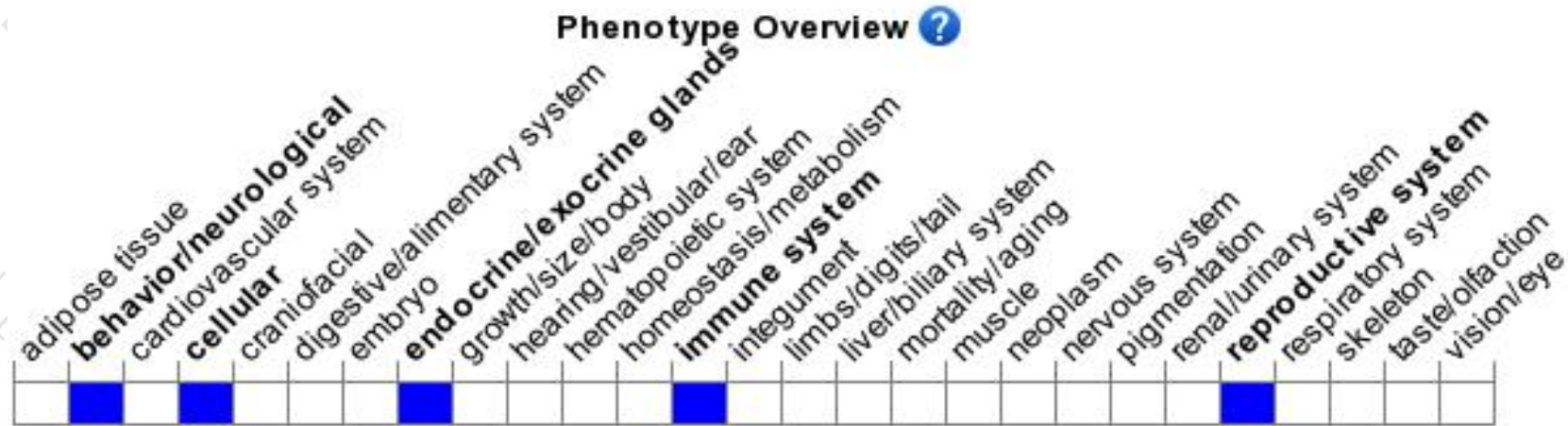
Genomic location distribution



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.

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