

Dolary Skock Co. Junb Cas9-CKO Strategy To hope of the color of the col

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Project Overview



Project Name

Junb

Project type

Cas9-CKO

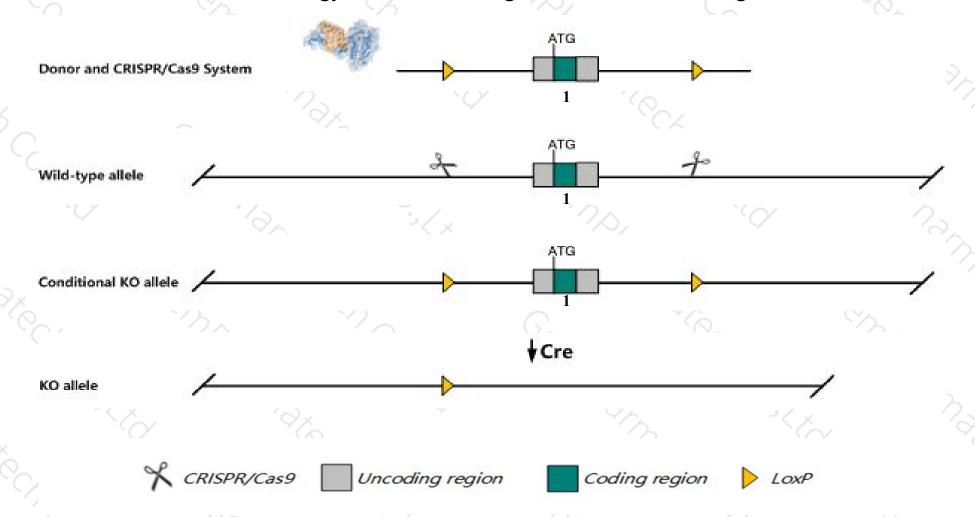
Strain background

C57BL/6JGpt

Conditional Knockout strategy



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



Technical routes



- The *Junb* gene has 2 transcripts. According to the structure of *Junb* gene, exon1 of *Junb-201* (ENSMUST00000064922.6) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Junb* 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, Homozygous null mutants die between embryonic day 8.5-10 due to impaired placental development. Embryos are severely growth retarded, but cell proliferation is normal, reflecting a failure to establish vascular interactions with the maternal circulation.
- > The *Junb* gene is located on the Chr8. 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)



Junb jun B proto-oncogene [Mus musculus (house mouse)]

Gene ID: 16477, updated on 19-Nov-2019

Summary

2 2

Official Symbol Junb provided by MGI

Official Full Name jun B proto-oncogene provided by MGI

Primary source MGI:MGI:96647

See related Ensembl: ENSMUSG00000052837

Gene type protein coding
RefSeq status VALIDATED
Organism <u>Mus musculus</u>

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

Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

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
Junb-201	ENSMUST00000064922.6	1809	344aa	Protein coding	CCDS22488	P09450 Q569U6	TSL:NA GENCODE basic APPRIS P1
Junb-202	ENSMUST00000209372.1	492	No protein	IncRNA		5	TSL:1

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

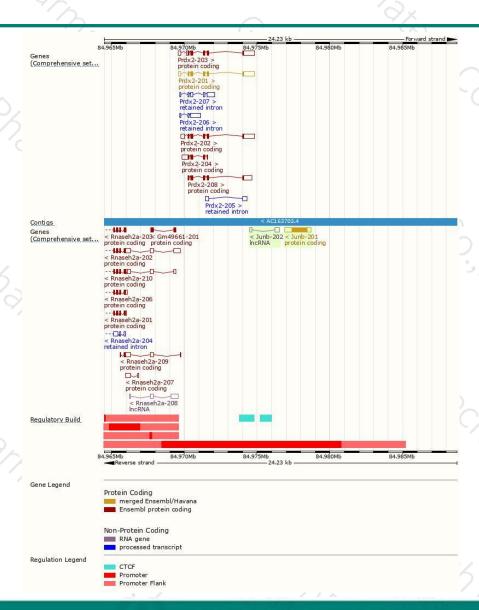
< Junb-201 protein coding

Reverse strand

1.81 kb

Genomic location distribution





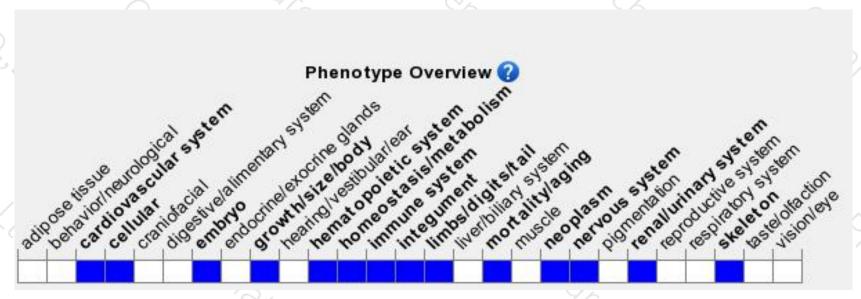
Protein domain





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, Homozygous null mutants die between embryonic day 8.5-10 due to impaired placental development. Embryos are severely growth retarded, but cell proliferation is normal, reflecting a failure to establish vascular interactions with the maternal circulation.



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





