

Mnx1 Cas9-CKO Strategy

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Reviewer:

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Design Date:

2019-12-16

Project Overview



Project Name

Mnx1

Project type

Cas9-CKO

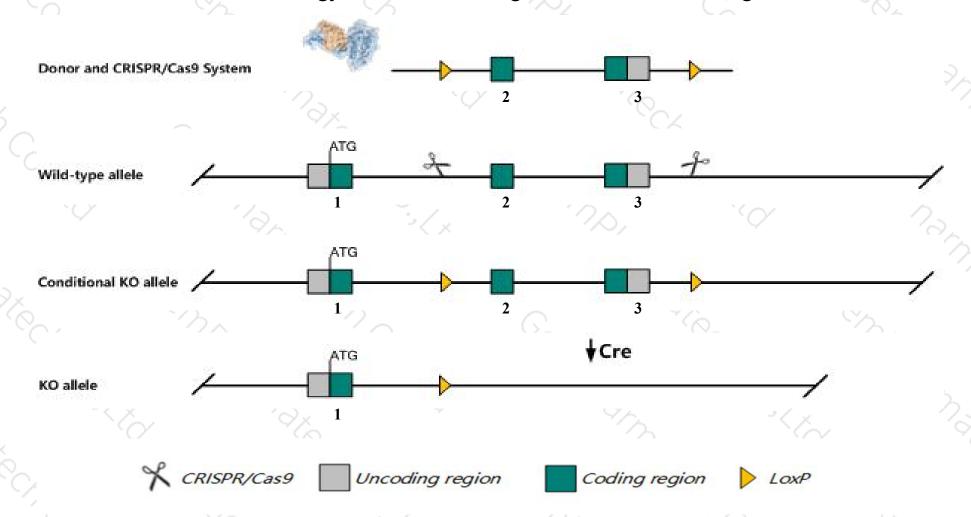
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Mnx1* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Mnx1* gene has 2 transcripts. According to the structure of *Mnx1* gene, exon2-exon3 of *Mnx1-202* (ENSMUST00000165512.3) transcript is recommended as the knockout region. The region contains 524bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Mnx1* 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 mice die at birth exhibiting pancreas dorsal lobe agenesis, small pancreatic islets, and aberrant beta-cell function and motor axon guidance. Mice homozygous for other reporter/null alleles show neonatal death, atelectasis, and impaired motor neuron and pancreas differentiation.
- The *Mnx1* gene is located on the Chr5. 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)



Mnx1 motor neuron and pancreas homeobox 1 [Mus musculus (house mouse)]

Gene ID: 15285, updated on 2-Apr-2019

Summary

☆ ?

Official Symbol Mnx1 provided by MGI

Official Full Name motor neuron and pancreas homeobox 1 provided by MGI

Primary source MGI:MGI:109160

See related Ensembl:ENSMUSG00000001566

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 HB9, HIxb9, MNR2

Expression Biased expression in duodenum adult (RPKM 2.3), colon adult (RPKM 2.0) and 5 other tissuesSee more

Orthologs <u>human all</u>

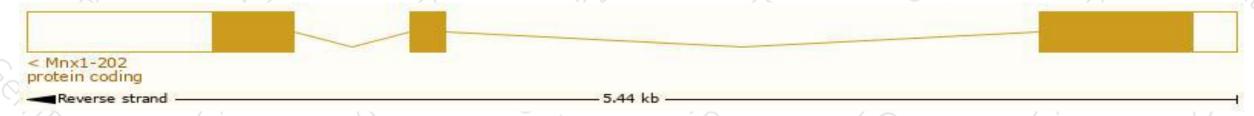
Transcript information (Ensembl)



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

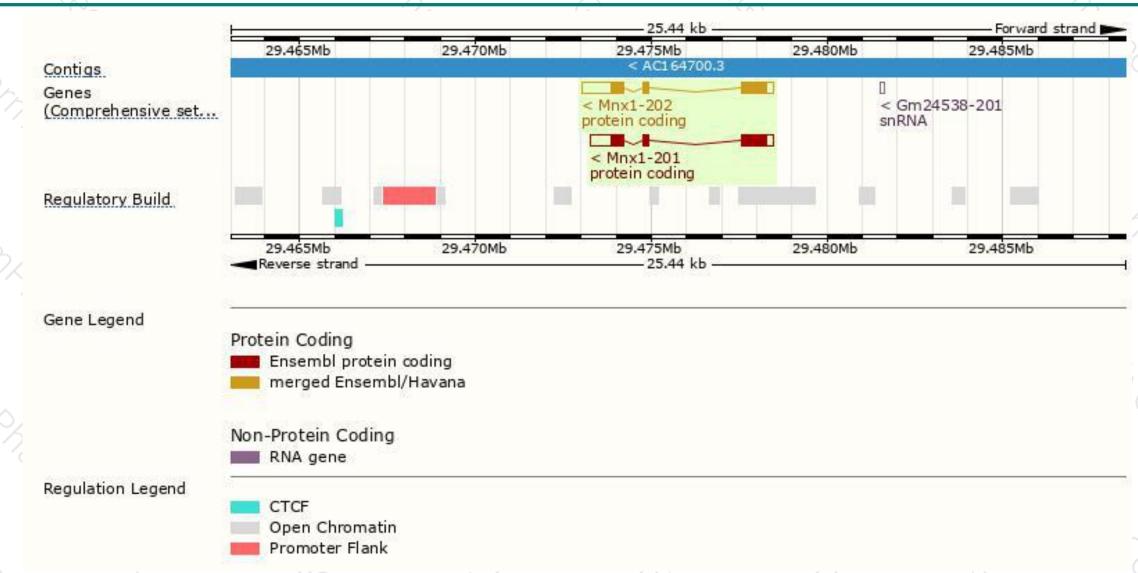
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Mnx1-202	ENSMUST00000165512.3	2245	404aa	Protein coding	CCDS19150	A2RSX2	TSL:1 GENCODE basic APPRIS P2
Mnx1-201	ENSMUST00000001608.7	2018	<u>404aa</u>	Protein coding	. 8	Q9QZW9	TSL:5 GENCODE basic APPRIS ALT2

The strategy is based on the design of Mnx1-202 transcript, The transcription is shown below



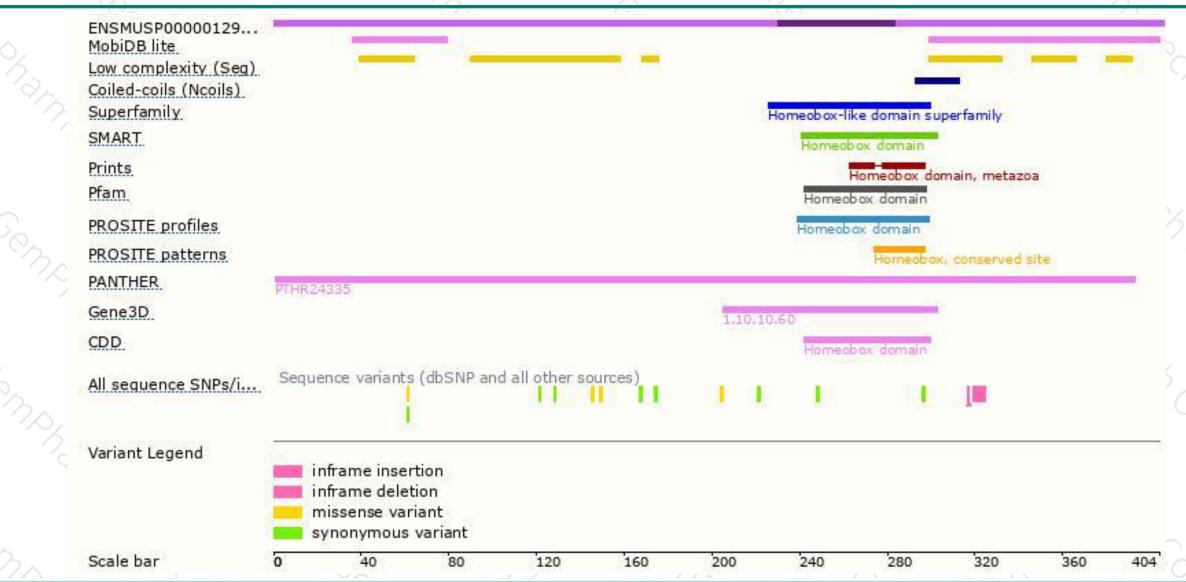
Genomic location distribution





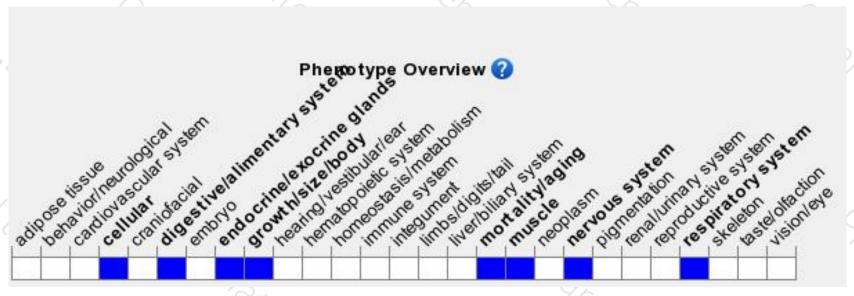
Protein domain





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, Homozygous null mice die at birth exhibiting pancreas dorsal lobe agenesis, small pancreatic islets, and aberrant beta-cell function and motor axon guidance. Mice homozygous for other reporter/null alleles show neonatal death, atelectasis, and impaired motor neuron and pancreas differentiation.



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





