

Hoxc9 Cas9-CKO Strategy

Designer:

Reviewer:

Design Date:

Yang Zeng

Xueting Zhang

2019-11-29

Project Overview



Project Name

Hoxc9

Project type

Cas9-CKO

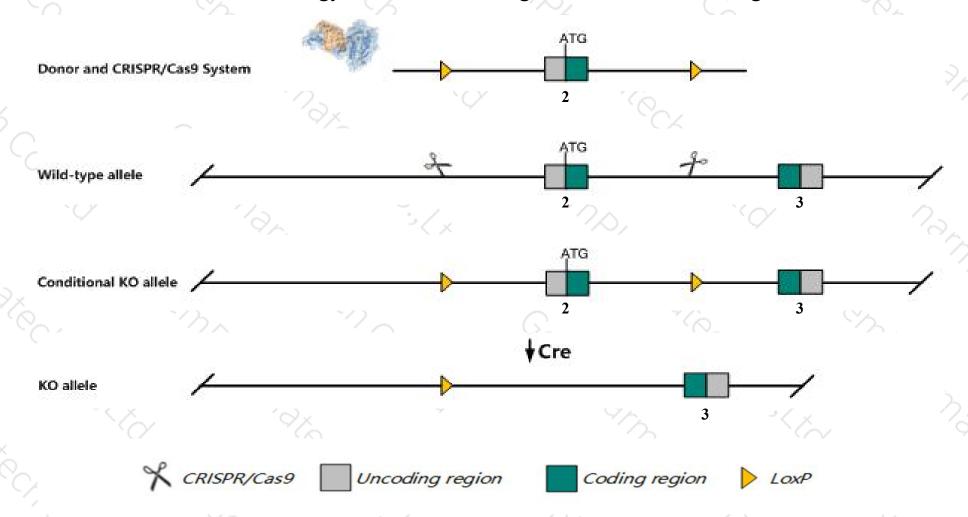
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Hoxc9* gene. The schematic diagram is as follows:



Technical routes



- The *Hoxc9* gene has 2 transcripts. According to the structure of *Hoxc9* gene, exon2 of *Hoxc9-201* (ENSMUST0000001706.6) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Hoxc9* 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 disruptions in this gene grow more slowly than normal and develop hunched backs. Forward transformations seen in vertebrae from L1 and forward to around T10. Abnormalities in the sternum and ribs attachments to the sternum are also seen.
- The Floxed region overlaps with intron1 of *Hoxc5-202* transcript. Knockout the region may affect the function of *Hoxc5-202* transcript.
- > The *Hoxc9* gene is located on the Chr15. 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)



Hoxc9 homeobox C9 [Mus musculus (house mouse)]

Gene ID: 15427, updated on 12-Nov-2019

Summary



Official Symbol Hoxc9 provided by MGI

Official Full Name homeobox C9 provided by MGI

Primary source MGI:MGI:96199

See related Ensembl:ENSMUSG00000036139

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 Hox-3.2

Expression Biased expression in ovary adult (RPKM 7.9), mammary gland adult (RPKM 7.9) and 7 other tissues See more

Orthologs human all



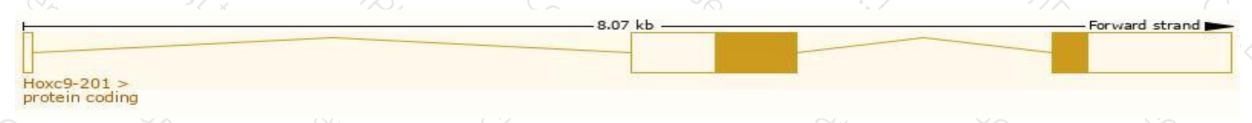
Transcript information (Ensembl)



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

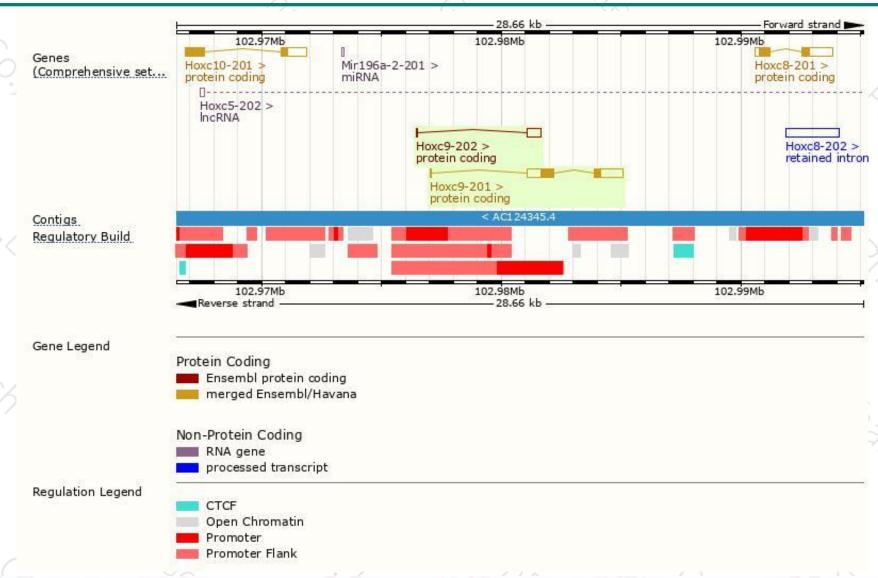
Name 🍦	Transcript ID 🍦	bp 🛊	Protein	Translation ID	Biotype 🍦	CCDS .	UniProt 4	Flags		
Hoxc9-201	ENSMUST00000001706.6	2374	<u>260aa</u>	ENSMUSP00000001706.6	Protein coding	CCDS27893₽	<u>P09633</u> ₽	TSL:1	GENCODE basic	APPRIS P1
Hoxc9-202	ENSMUST00000173306.1	634	<u>5aa</u>	ENSMUSP00000134197.1	Protein coding	(4-3)	1-		CDS 3' incomplete	TSL:3

The strategy is based on the design of *Hoxc9-201* transcript, The transcription is shown below



Genomic location distribution





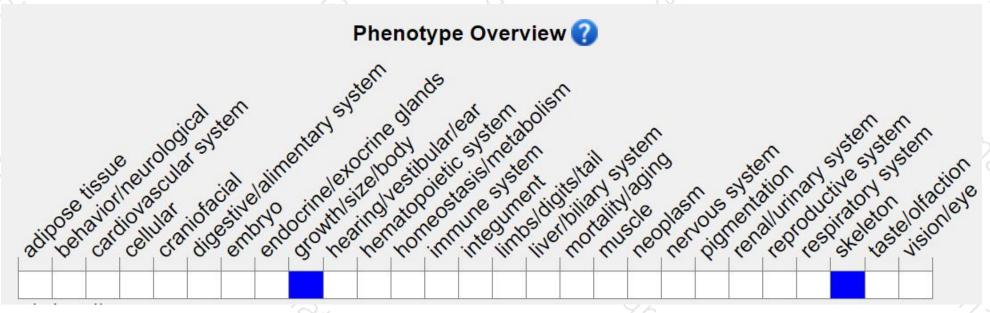
Protein domain





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 disruptions in this gene grow more slowly than normal and develop hunched backs. Forward transformations seen in vertebrae from L1 and forward to around T10. Abnormalities in the sternum and ribs attachments to the sternum are also seen.



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





