

Irx1 Cas9-CKO Strategy

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Design Date: 2019-7-25

Project Overview



Project Name

Irx1

Project type

Cas9-CKO

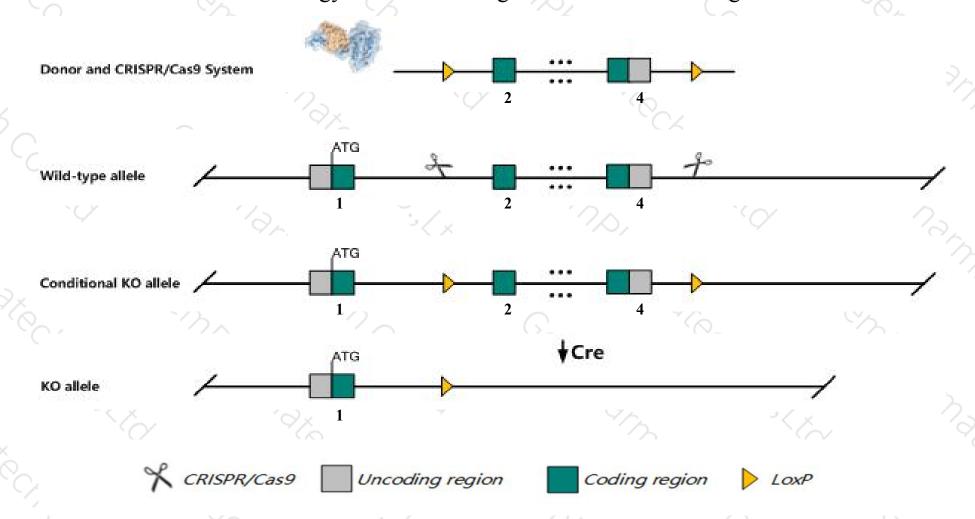
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Irx1* gene. The schematic diagram is as follows:



Technical routes



- The *Irx1* gene has 3 transcripts. According to the structure of *Irx1* gene, exon2-exon4 of *Irx1-201* (ENSMUST00000077337.8) transcript is recommended as the knockout region. The region contains most 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 *Irx1* 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 a knock-out allele exhibit complete neonatal lethality associated with delayed lung maturation, cyanosis, respiratory distress, atelectasis and decreased surfactant secretion, and show defects in dental epithelial cell differentiation and delayed growth of the lower incisors.
- The *Irx1* gene is located on the Chr13. 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)



Irx1 Iroquois homeobox 1 [Mus musculus (house mouse)]

Gene ID: 16371, updated on 31-Jan-2019

Summary

△ ?

Official Symbol Irx1 provided by MGI

Official Full Name Iroquois homeobox 1 provided by MGI

Primary source MGI:MGI:1197515

See related Ensembl:ENSMUSG00000060969

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

Expression Biased expression in lung adult (RPKM 102.7), mammary gland adult (RPKM 39.1) and 8 other tissues See more

Orthologs <u>human</u> all

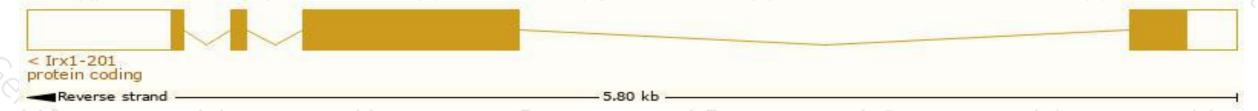
Transcript information (Ensembl)



The gene has 3 transcripts, all transcripts are shown below:

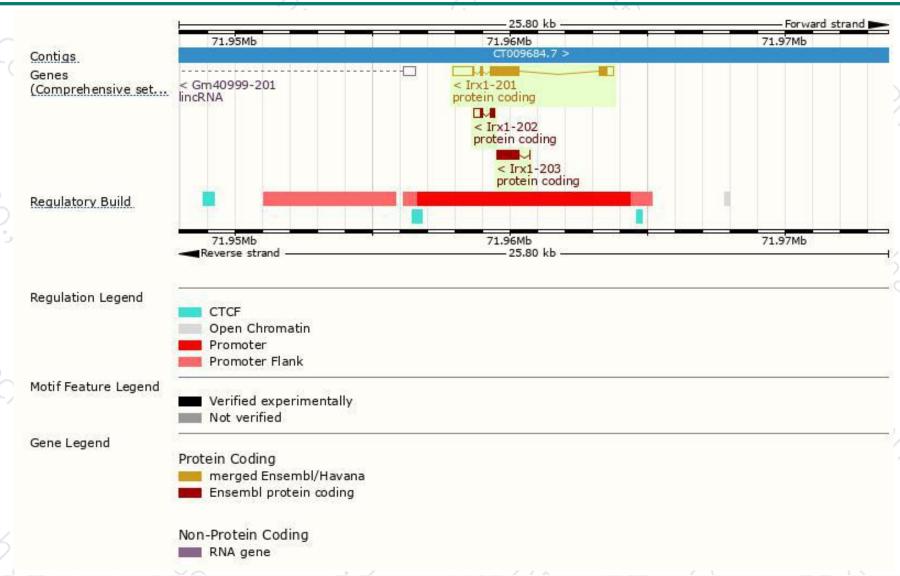
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Irx1-201	ENSMUST00000077337.8	2373	480aa	Protein coding	CCDS26627	P81068	TSL:1 GENCODE basic APPRIS P1
Irx1-203	ENSMUST00000223460.1	820	273aa	Protein coding	* .	A0A1Y7VIS3	5' and 3' truncations in transcript evidence prevent annotation of the start and the end of the CDS. CDS 5' and 3' incomplete TSL:3
Irx1-202	ENSMUST00000223379.1	498	86aa	Protein coding	-	A0A1Y7VJ15	CDS 5' incomplete TSL:2

The strategy is based on the design of *Irx1-201* transcript, The transcription is shown below



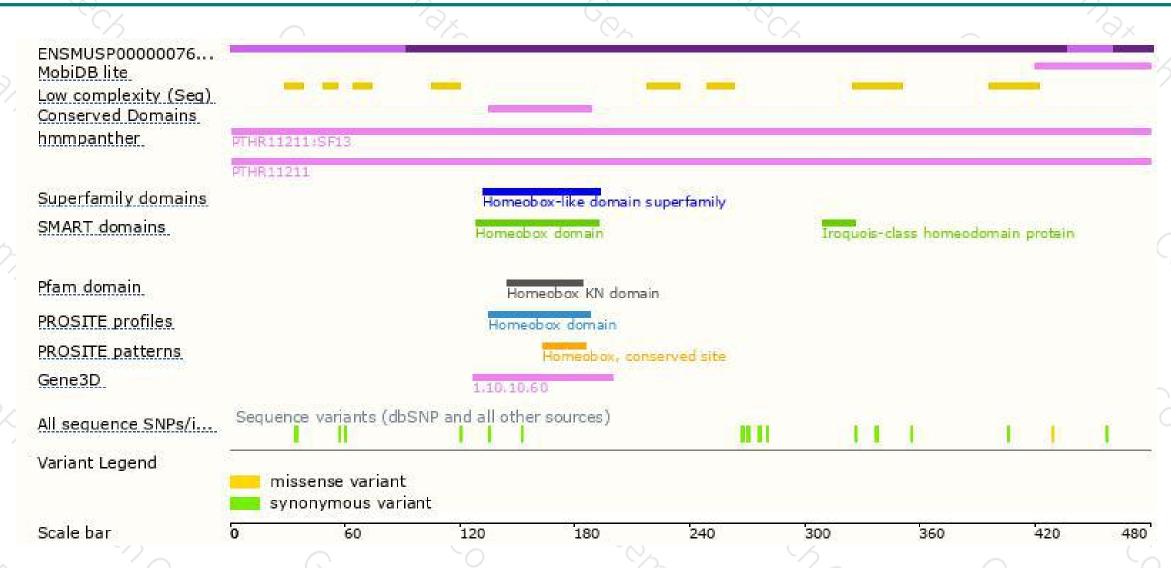
Genomic location distribution





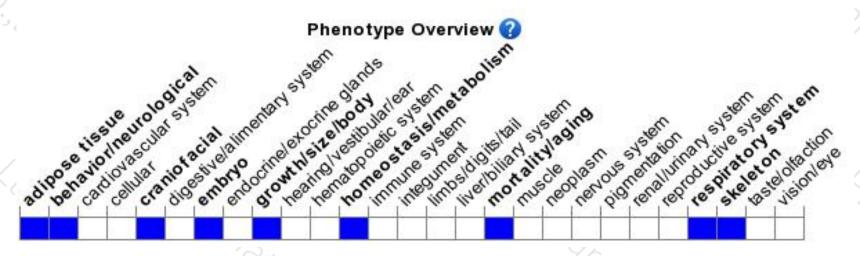
Protein domain





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 a knock-out allele exhibit complete neonatal lethality associated with delayed lung maturation, cyanosis, respiratory distress, at electasis and decreased surfactant secretion, and show defects in dental epithelial cell differentiation and delayed growth of the lower incisors.



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





