

***Nkx1-1* Cas9-KO Strategy**

Designer: JiaYu

Reviewer: Xiaojing Li

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Project Overview

Project Name

Nkx1-1

Project type

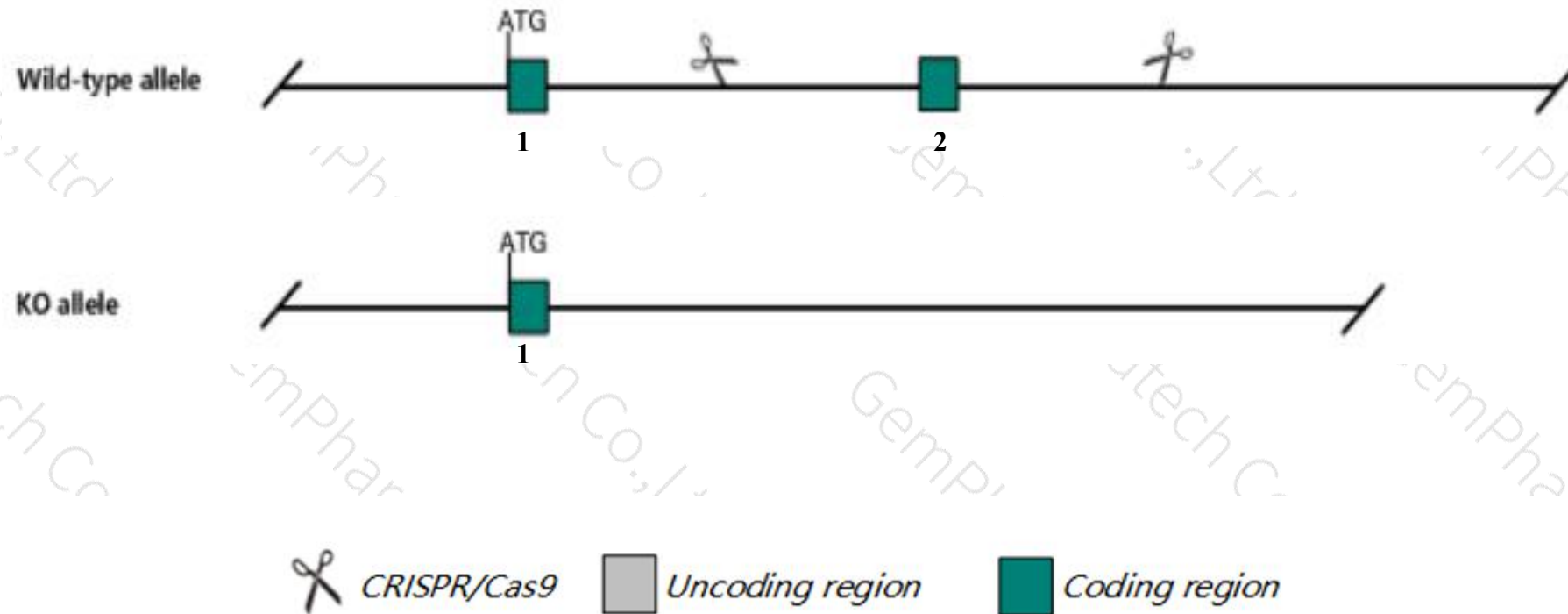
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Nkx1-1* gene. The schematic diagram is as follows:



- The *Nkx1-1* gene has 1 transcript. According to the structure of *Nkx1-1* gene, exon2 of *Nkx1-1-201*(ENSMUST00000173348.1) transcript is recommended as the knockout region. The region contains 857bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Nkx1-1* gene. The brief process is as follows: CRISPR/Cas9 system 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.

- According to the existing MGI data, mice homozygous for disruptions in this gene show poor growth and survival. Most die within the first three weeks of life. Those that reach adulthood are fertile but do not produce viable offspring.
- The *Nkx1-1* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

Nkx1-1 NK1 homeobox 1 [Mus musculus (house mouse)]

Gene ID: 672284, updated on 13-Mar-2020

Summary



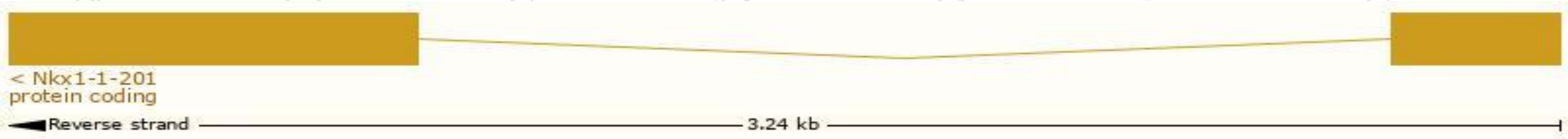
Official Symbol	Nkx1-1 provided by MGI
Official Full Name	NK1 homeobox 1 provided by MGI
Primary source	MGI:MGI:109346
See related	Ensembl:ENSMUSG00000029112
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Nkx-1.1, Sax2
Expression	Low expression observed in reference dataset See more
Orthologs	human all

Transcript information (Ensembl)

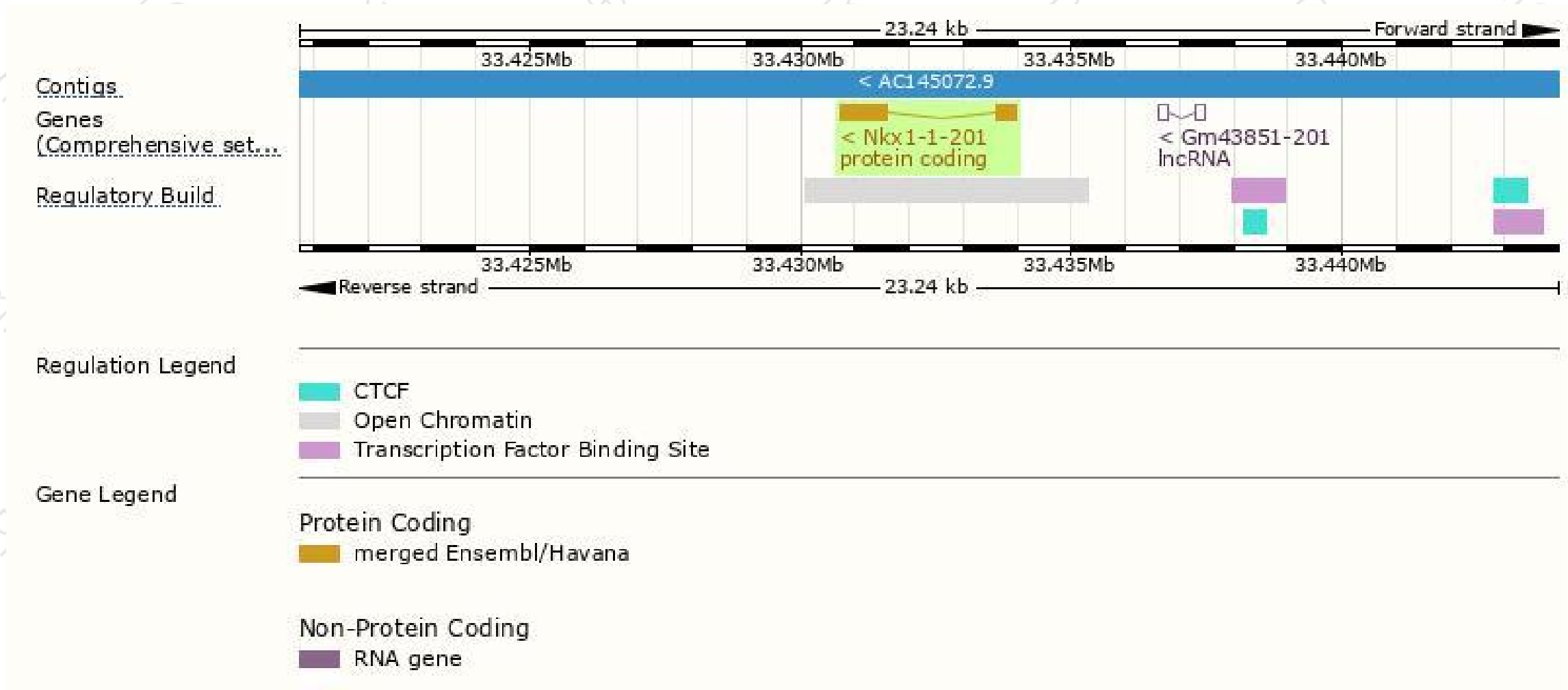
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Nkx1-1-201	ENSMUST00000173348.1	1209	402aa	Protein coding	-	G3UXB3	TSL:5 GENCODE basic APPRIS P1

The strategy is based on the design of *Nkx1-1-201* transcript, the transcription is shown below:



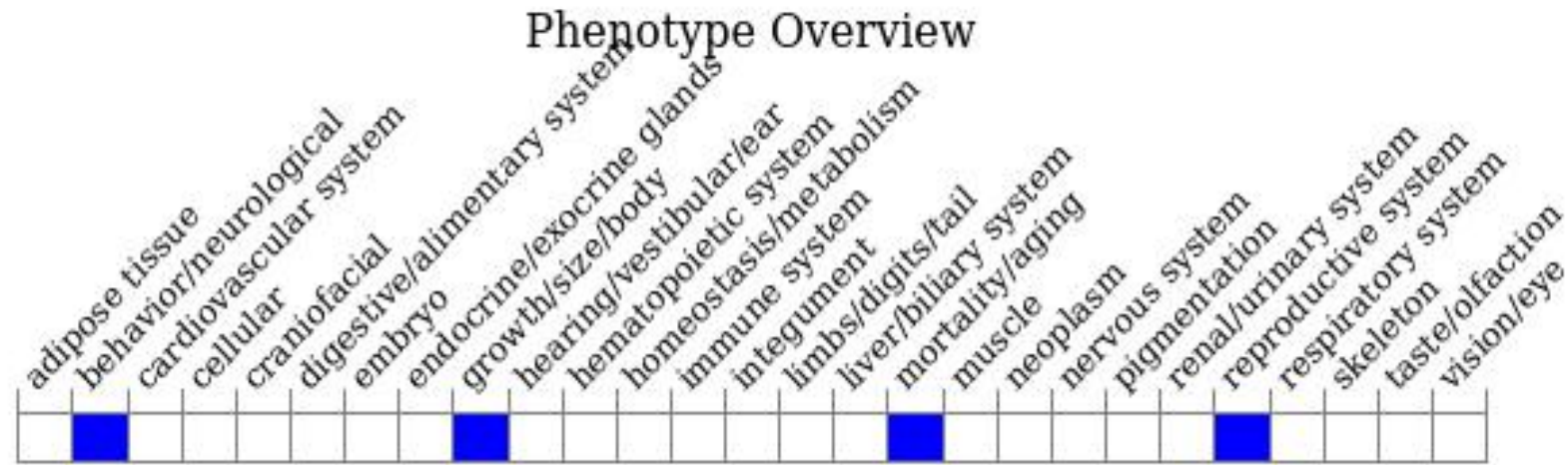
Genomic location distribution



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 show poor growth and survival.

Most die within the first three weeks of life. Those that reach adulthood are fertile but do not produce viable offspring.

If you have any questions, you are welcome to inquire.

Tel: 400-9660890

