

# ***Lhx5*** Cas9-KO Strategy

**Designer: Xueting Zhang**

**Reviewer: Daohua Xu**

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# Project Overview

**Project Name**

***Lhx5***

**Project type**

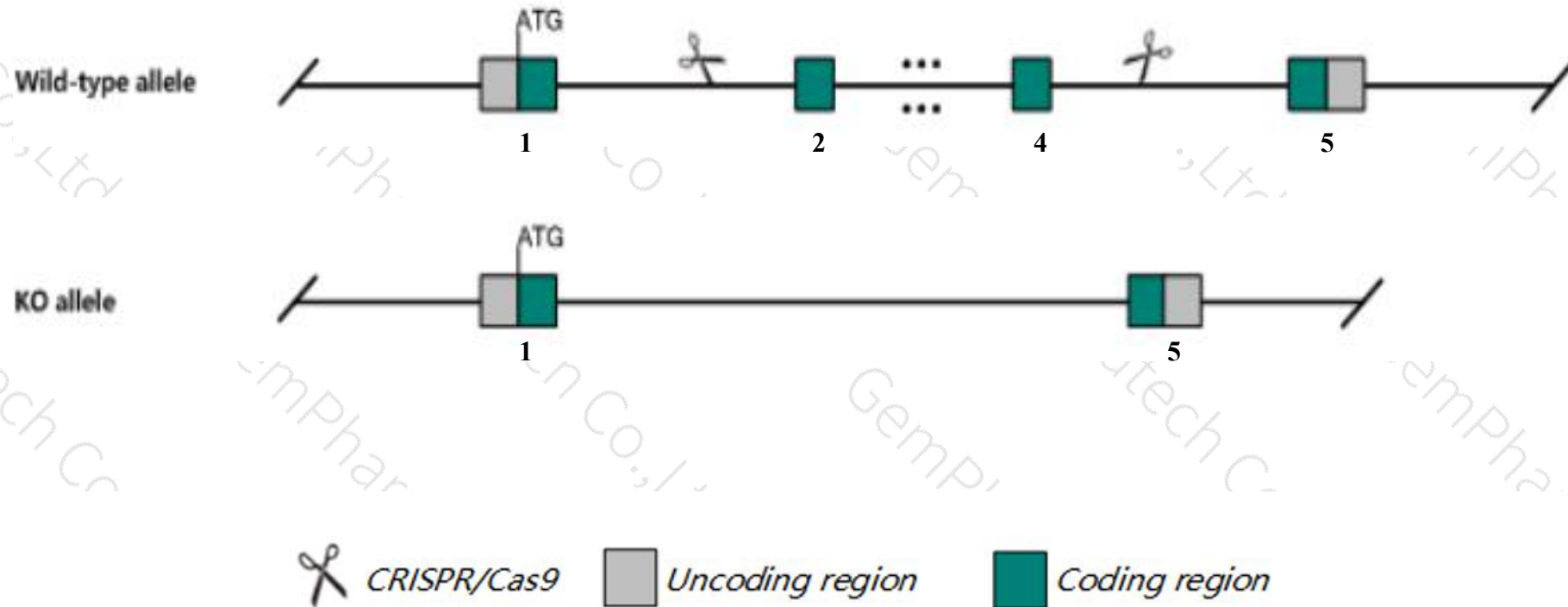
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Lhx5* gene. The schematic diagram is as follows:



- The *Lhx5* gene has 1 transcript. According to the structure of *Lhx5* gene, exon2-exon4 of *Lhx5*-201(ENSMUST00000031591.9) transcript is recommended as the knockout region. The region contains 668bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Lhx5* 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, most mice homozygous for a null mutation display defective hippocampal development and die within a few days after birth. Postmitotic hippocampal cells are unable to differentiate properly and migrate to correct positions, resulting in structural anomalies of the Ammon's horn and the dentate gyrus.
- The knockout region is near to the N-terminal of *Gm27199* gene, this strategy may influence the regulatory function of the N-terminal of *Gm27199* gene.
- The *Lhx5* 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)

Lhx5 LIM homeobox protein 5 [Mus musculus (house mouse)]

Gene ID: 16873, updated on 13-Mar-2020

## Summary



Official Symbol Lhx5 provided by [MGI](#)

Official Full Name LIM homeobox protein 5 provided by [MGI](#)

Primary source [MGI:MGI:187792](#)

See related [Ensembl:ENSMUSG00000029595](#)

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 Lim2

Expression Biased expression in CNS E11.5 (RPKM 6.4), whole brain E14.5 (RPKM 6.0) and 5 other tissues [See more](#)

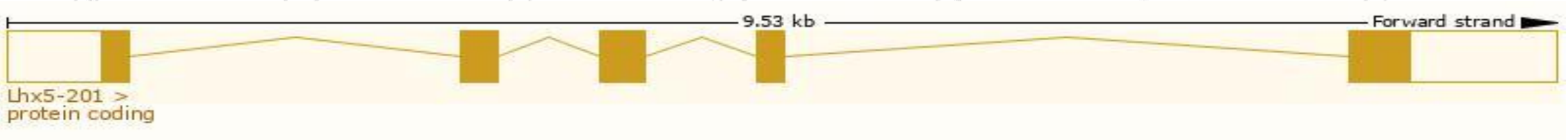
Orthologs [human all](#)

# Transcript information (Ensembl)

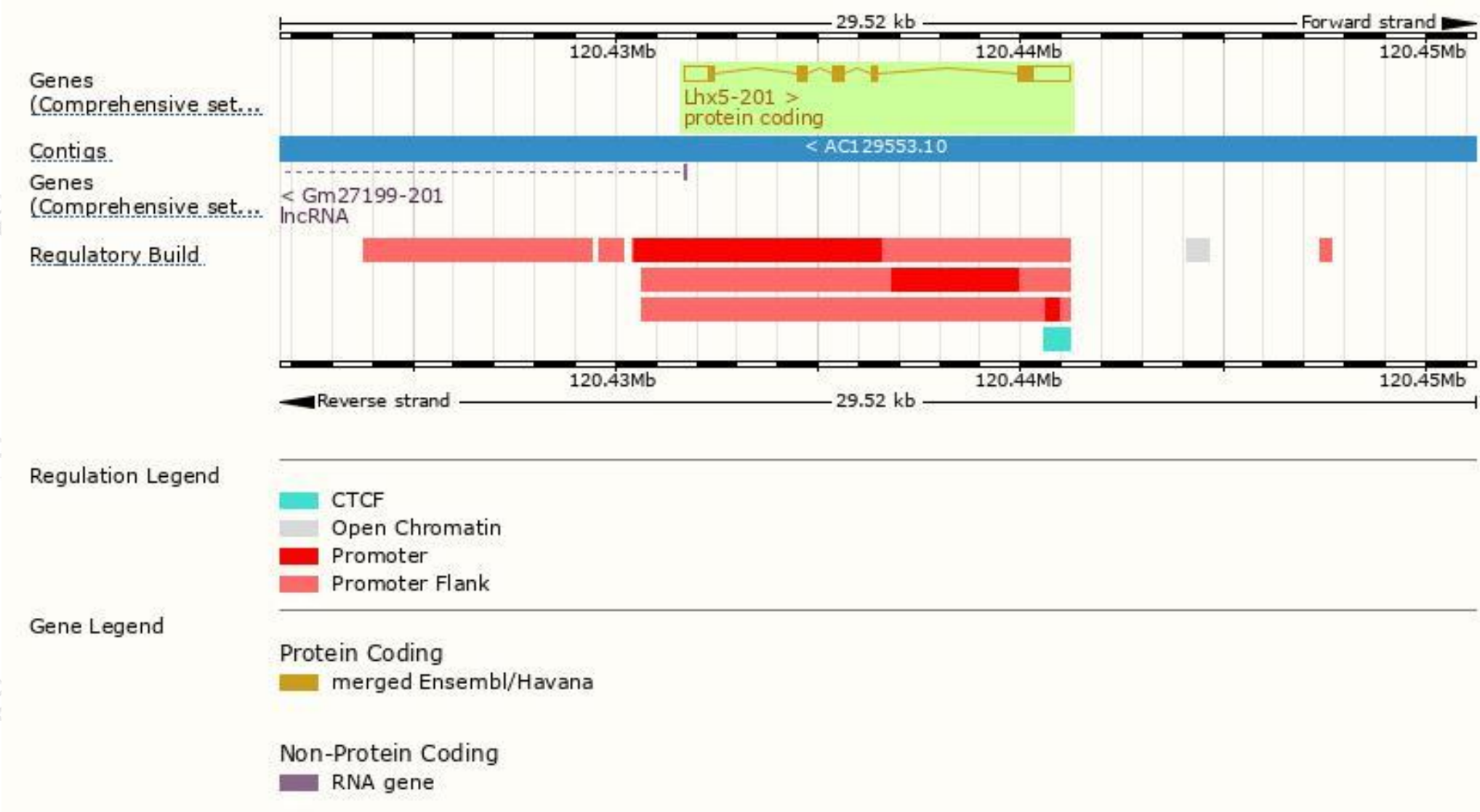
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Lhx5-201	<a href="#">ENSMUST00000031591.9</a>	2689	<a href="#">402aa</a>	Protein coding	<a href="#">CCDS19617</a>	<a href="#">P61375 Q543P4</a>	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Lhx5-201* transcript, the transcription is shown below:



# Genomic location distribution

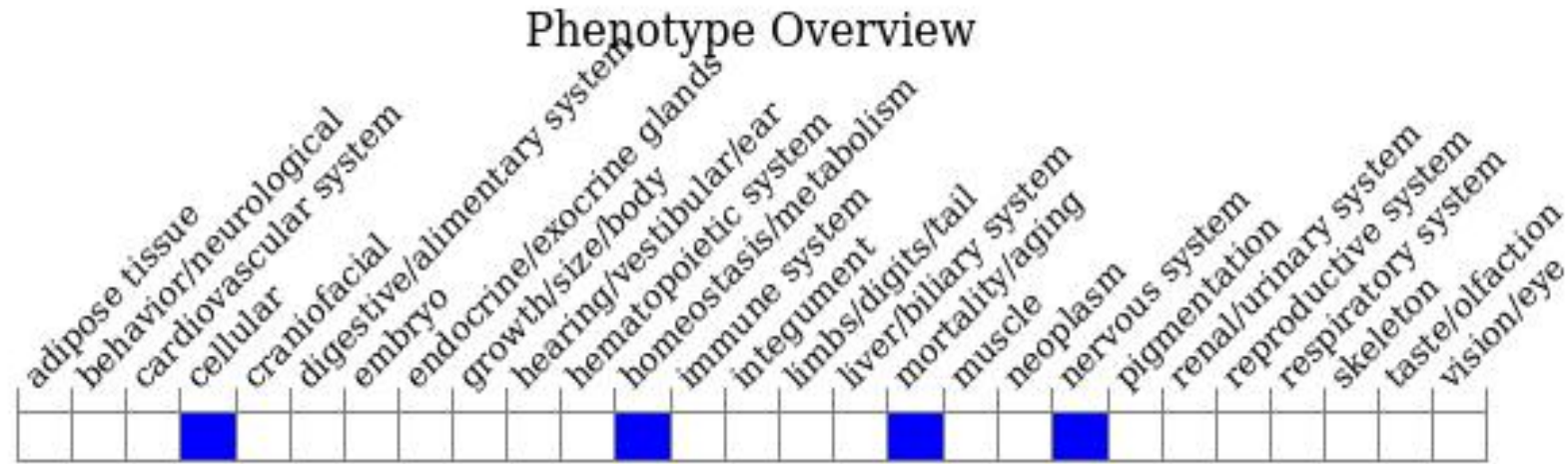




# Protein domain



# 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, most mice homozygous for a null mutation display defective hippocampal development and die within a few days after birth. Postmitotic hippocampal cells are unable to differentiate properly and migrate to correct positions, resulting in structural anomalies of the Ammon's horn and the dentate gyrus.

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

Tel: 400-9660890

