

Lhx5 Cas9-CKO Strategy

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Project Overview

Project Name

Lhx5

Project type

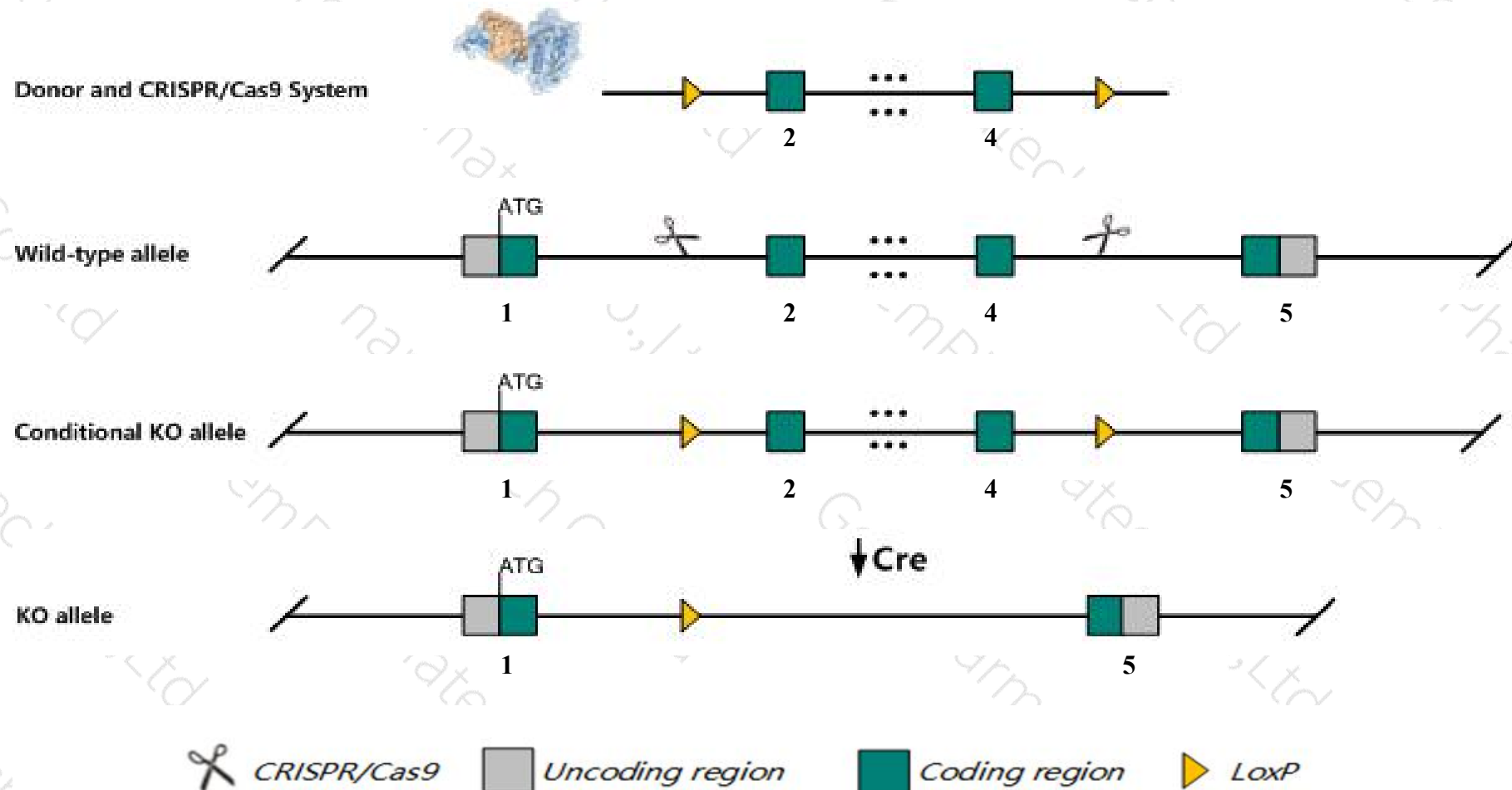
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

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



Technical routes

- 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 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.

- 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 floxed 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological 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](#)

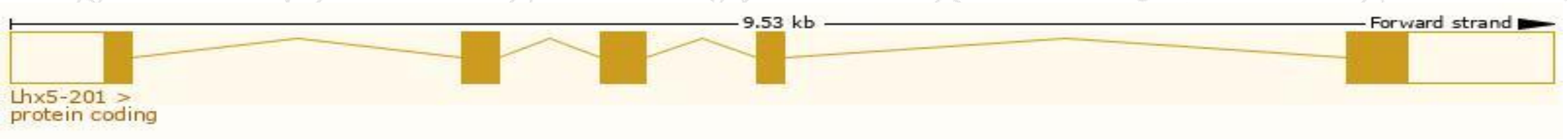
Orthologs [human all](#)

Transcript information (Ensembl)

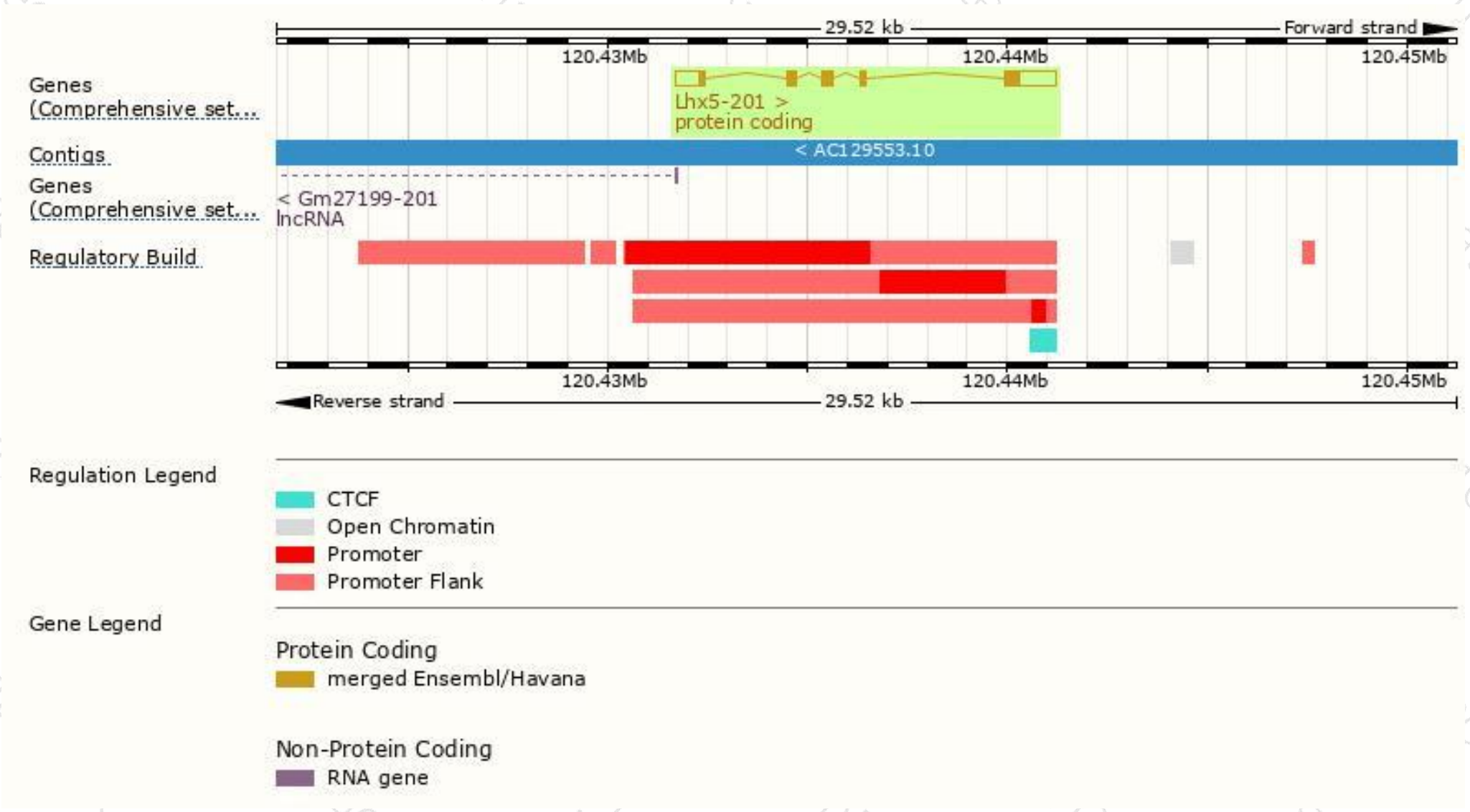
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Lhx5-201	ENSMUST00000031591.9	2689	402aa	Protein coding	CCDS19617	P61375 Q543P4	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

ENSMUSP00000031...

MobiDB lite

Low complexity (Seq)

Superfamily

SMART

Pfam

PROSITE profiles

PROSITE patterns

PANTHER

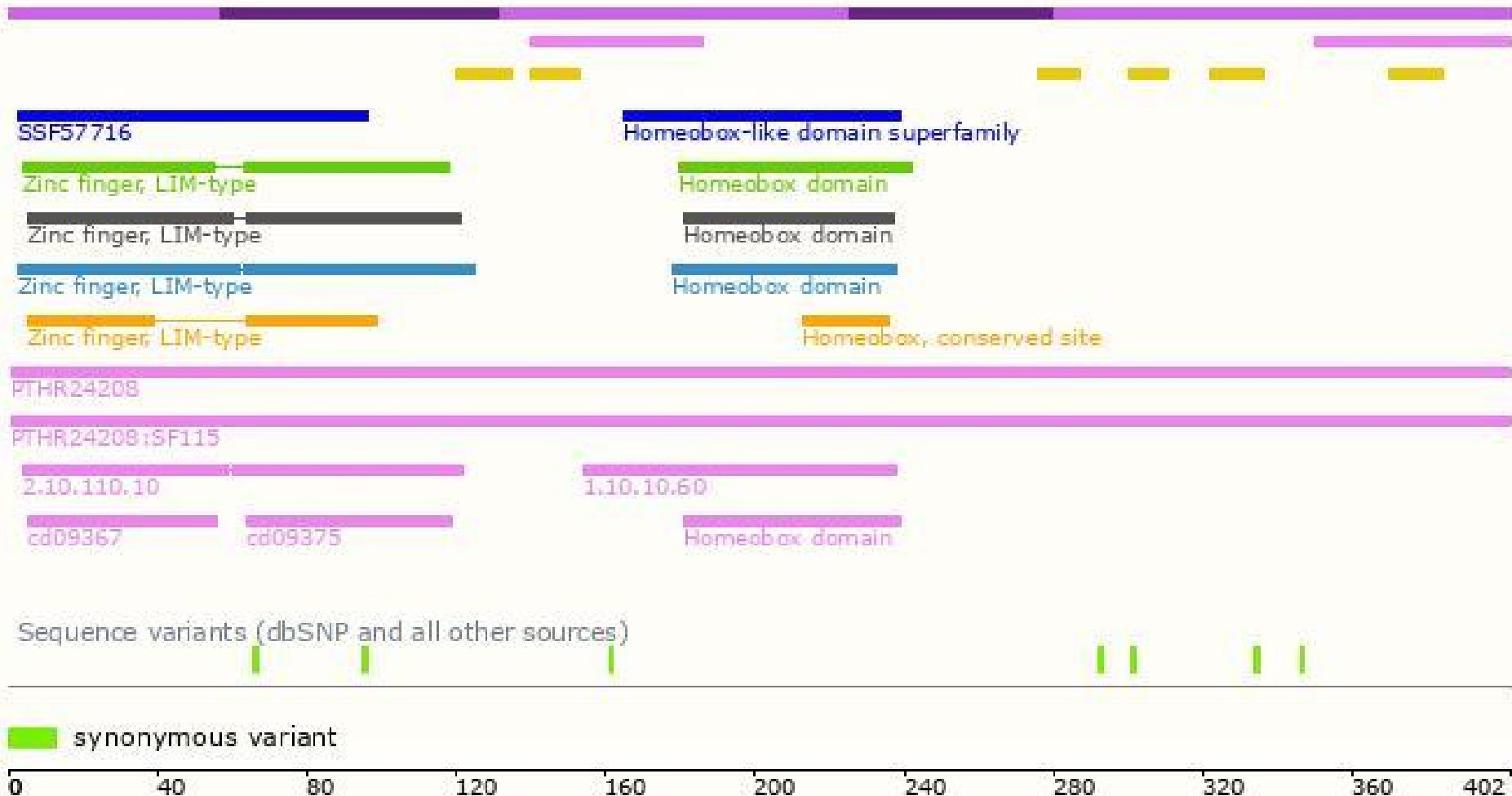
Gene3D

CDD

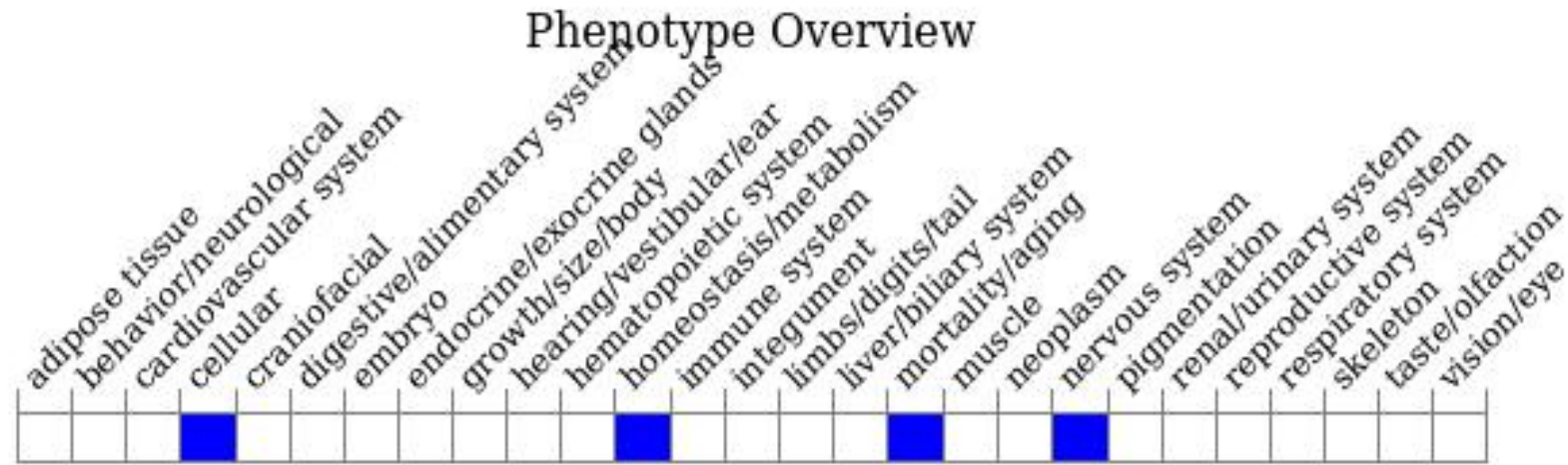
All sequence SNPs/i...

Variant Legend

Scale bar



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.

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