

***Chrna2* Cas9-KO Strategy**

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Project Overview

Project Name

Chrna2

Project type

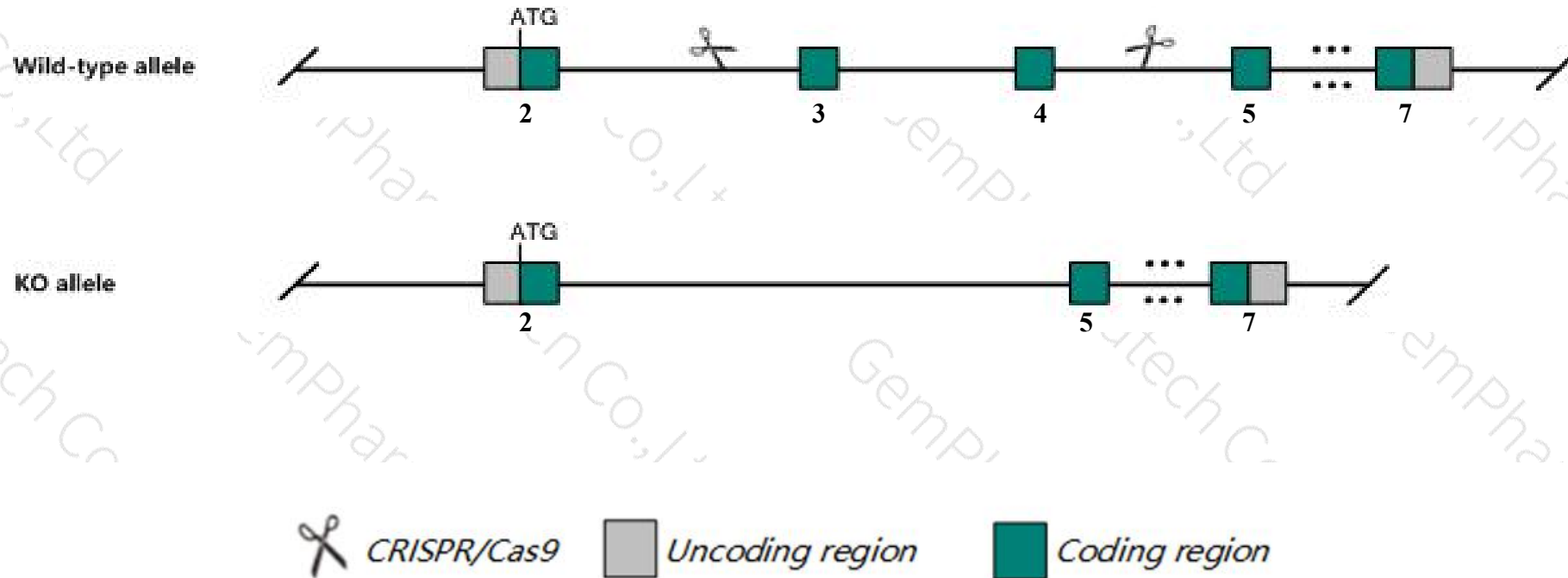
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Chrna2* gene. The schematic diagram is as follows:



- The *Chrna2* gene has 2 transcripts. According to the structure of *Chrna2* gene, exon3-exon4 of *Chrna2-201* (ENSMUST00000022620.10) transcript is recommended as the knockout region. The region contains 197bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Chrna2* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a targeted null mutation do not exhibit any significant abnormalities compared to controls.
- The *Chrna2* gene is located on the Chr14. 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)

Chrna2 cholinergic receptor, nicotinic, alpha polypeptide 2 (neuronal) [Mus musculus (house mouse)]

Gene ID: 110902, updated on 31-Jan-2019

Summary



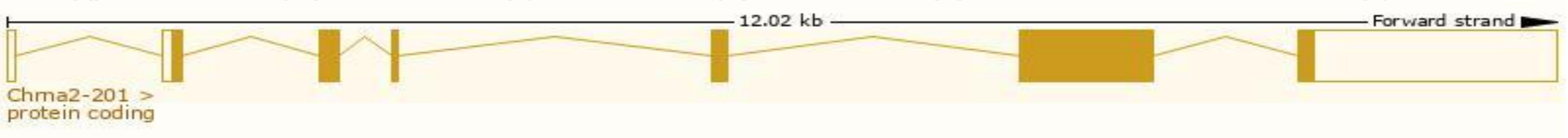
Official Symbol	Chrna2 provided by MGI
Official Full Name	cholinergic receptor, nicotinic, alpha polypeptide 2 (neuronal) provided by MGI
Primary source	MGI:MGI:87886
See related	Ensembl:ENSMUSG00000022041
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Acra-2, Acra2, BC011490
Expression	Biased expression in genital fat pad adult (RPKM 6.8), ovary adult (RPKM 1.4) and 6 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

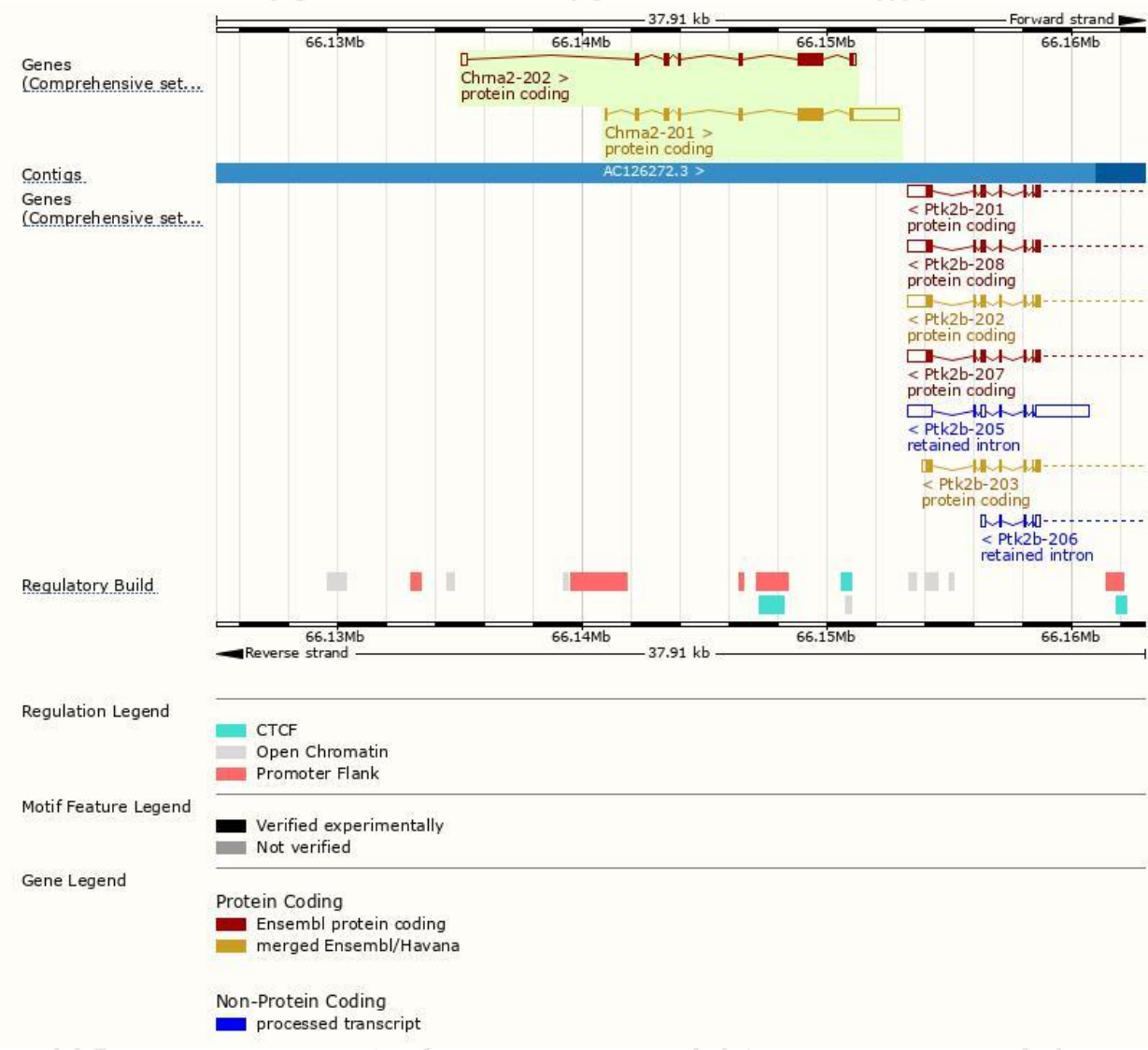
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Chrna2-201	ENSMUST00000022620.10	3554	512aa	Protein coding	CCDS27219	Q91X60	TSL:1 GENCODE basic APPRIS P1
Chrna2-202	ENSMUST00000206455.1	1933	512aa	Protein coding	CCDS27219	Q91X60	TSL:5 GENCODE basic APPRIS P1

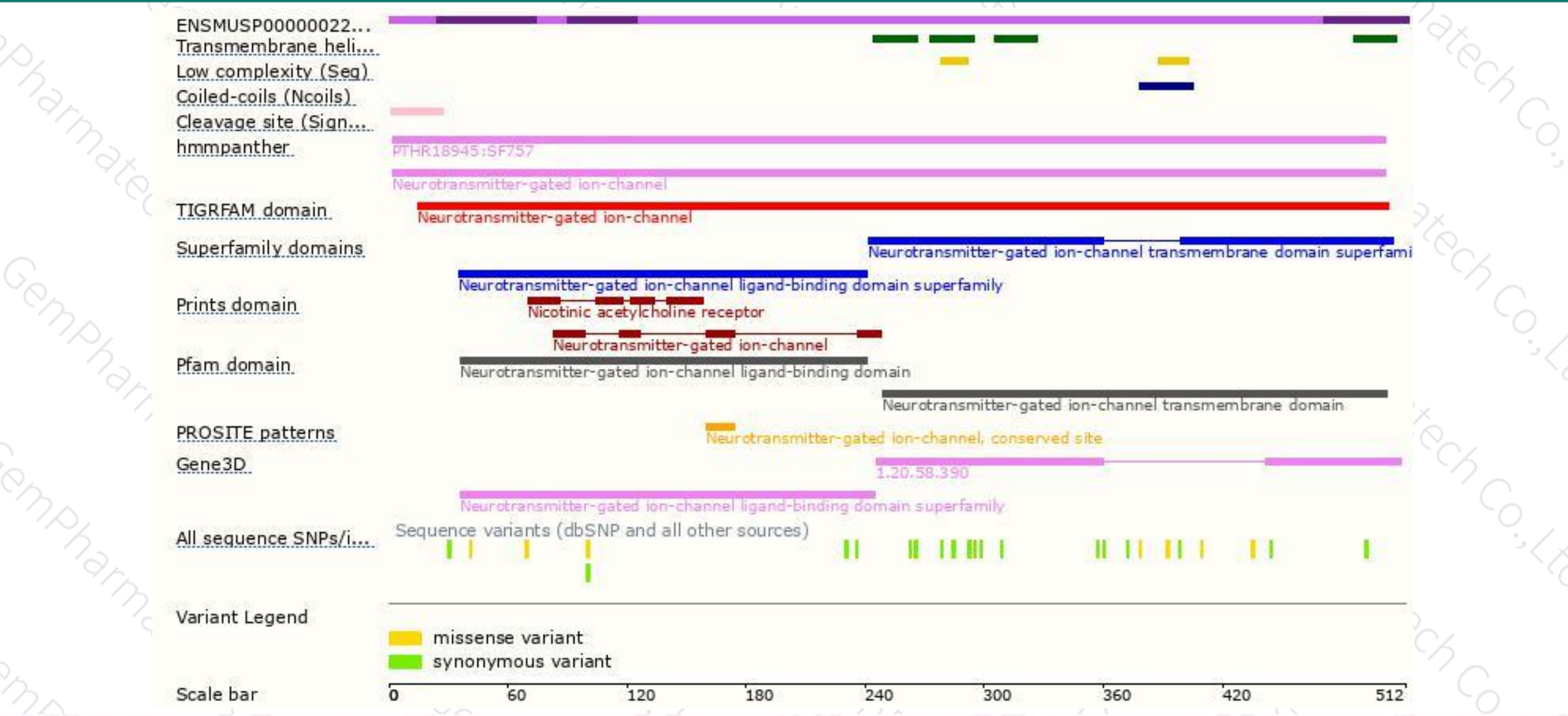
The strategy is based on the design of *Chrna2-201* transcript,The transcription is shown below



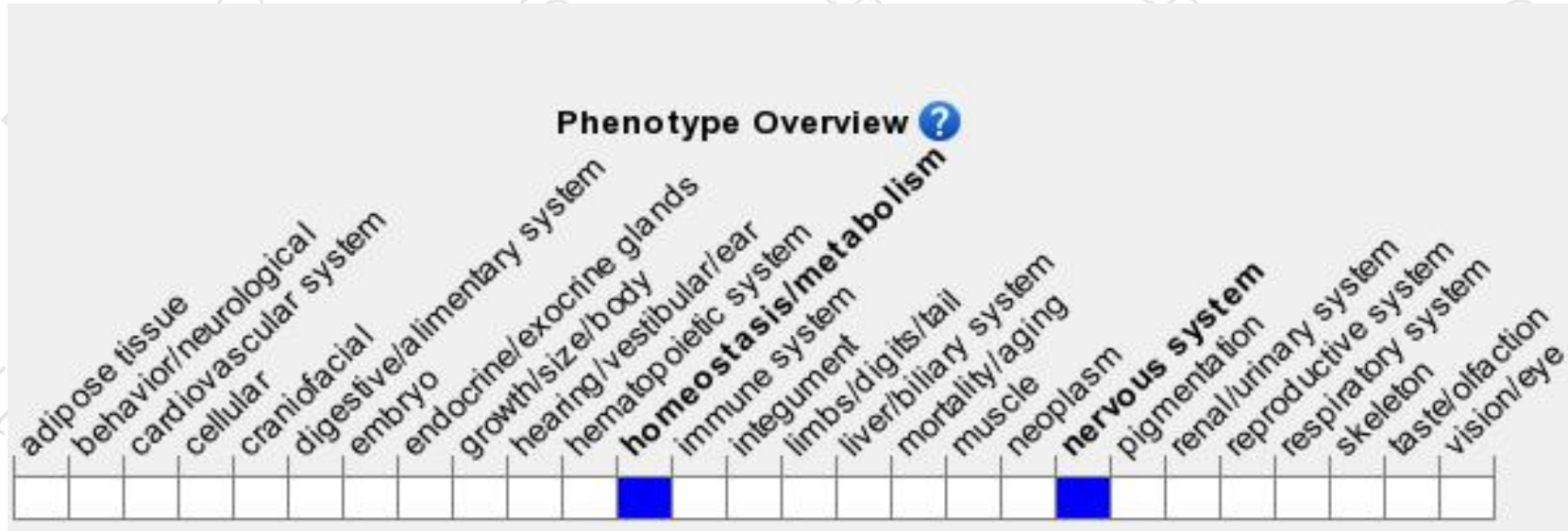
Genomic location distribution



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 targeted null mutation do not exhibit any significant abnormalities compared to controls.

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

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